U.S. DEPARTMENT OF COMMERCE Patent and Trademark Office

## SEARCH REQUEST FORM

equestor's ame:		Serial Number:	
ame:ame:	Dhone.		Art Unit:
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earch Topic: lease write a detailed statement of erms that may have a special mea lease attach a copy of the sequer	of search topic. Describe s aning. Give examples or r ice. You may include a co	specifically as possible the selevent citations, authors, keppy of the broadest and/or m	subject matter to be searched. Define any teywords, etc., if known. For sequences, tost relevent claim(s).
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Date completed:07-	07-04	Search Site	<b>Vendors</b> IG
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Elapsed time:		Type of Search	APS
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Number of Searches:		Structure	DARC/Questel
Number of Databases:		Bibliograp	ohic Other CGN

PTO-1590 (9-90)

From:

Ramirez, Delia

Sent:

Wednesday, June 30, 2004 11:45 AM STIC-Biotech/ChemLib

To:

Subject:

case 09/762,258

Hi,

I would like to request the following interference search:

- 1. seq id 1 and 2 in the nucleic acid databases
- 2. seq id 2 in the protein databases

Thank you,

Delia M. Ramirez, Ph.D. Patent Examiner Recombinant Enzymes-Art Unit 1652 USPTO 400 Dulany Street, Remsen Bldg., 3A74, Mail room 3C70 Alexandria, VA 22314 (571) 272-0938 delia.ramirez@uspto.gov

Searcher:	
Phone:	
Location:	
Date Picked Up:	
Date Completed:	
Searcher Prep/Review:	
Clerical:	
Online times	

TYPE OF SEARCH:
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other

VENDOR/COST (where applic.)
STN:
DIALOG:
Questel/Orbit:
DRLink:
Lexis/Nexis:
Sequence Sys.:
WWW/Internet:
Other (specify):

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Maximum Match 100%
Listing first 45 summaries
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     Query
Match
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1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*

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US-09-111-2709-5
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US-09-111-444-9
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   LPPYLTPDARDLVKKFLKRNPSQR1GGGPGDAADVQRHPFFRHMNWDDLLAWRVDPPFRP
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## ALIGNMENTS

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APPLICANT: Yohn Blenis
APPLICANT: Kay K. Lee-Fruman
APPLICANT: Calvin J. Kuo
APPLICANT: Calvin J. Kuo
TITLE OF INVENTION: P5456K AND P8586K GENES, PR
TITLE OF INVENTION: PRIMERS, PROBES, AND DETECT
FILE REFERENCE: 00246/506002
CURRENT APPLICATION NUMBER: US/09/430,564
CURRENT FILING DATE: 1999-10-29
PRIOR FILING DATE: 1999-10-29
PRIOR FILING DATE: 1998-10-29
APPLICATION NUMBER: 60/106,141
PRIOR FILING DATE: 1998-10-29
INUMBER OF SEQ ID NOS: 16
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 495
TYPE: PRT
COGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application Patent No. 6372467 GENERAL INFORMATION:
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301 LPPYLTPDARDLVKKFLKRNPSQRIGGGPGDAADVQRHPFFRHMWDDLLAWRVDPPFRP
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RESULT 2 US-09-430-564-2

GENERAL INFORMATION:

APPLICANT: APPLICANT: Š 밁 Ş 밁

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TITLE OF INVENTION: P5486K AND P8586K GENES, PROTEINS,
TITLE OF INVENTION: PRIMERS, PROBES, AND DETECTION METHODS
FILE REFERENCE: 00246/506002
CURRENT APPLICATION NUMBER: US/09/430,564
CURRENT FILING DATE: 1999-10-29
PRIOR APPLICATION NUMBER: 60/106,141
PRIOR FILING DATE: 1998-10-29
NUMBER: 005106,141
PRIOR FILING DATE: 1998-10-29
NUMBER: FASTSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 482
TYPE: PRT
ORGANISM: Homo sapiens
US-09-430-564-2
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                                             PVSPLKFSPFEGFRPSPSLPEPTELPLPPLLPPPPDSTTAPLPIRPPSGTKKSKRGRGRP
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Calvin J. Kuo
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Best Local (
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TOPOLOGY: linea
IMMEDIATE SOURCE:
LIBRARY: GenBan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Billings, Lucy J.
REGISTATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/7:
FILING DATE: Filed Herewith
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
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ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS,
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ATTORNEY/AGENT INFORMATION:
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APPLICANT: Goli, Surya K.
APPLICANT: Hillman, Jennifer L.
TITLE OF INVENTION: NOVEL HUVAN SERINE/THREONINE
TITLE OF INVENTION: PROTEIN KINASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Palo Alto
STATE: CA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
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5985635
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                                                                                                                                                                                                                                                AGLEP----VGHYBEVELTETSVNVGPERIGPHCFELLRVLGKGGYGKVFQVRKVQGTNL 105
  FGLCKESIHEGAVTHTFCGTIEYMAPEILVRSGHNRAVDWWSLGALMYDMLTGSPFFTAE
                                           GGELFMQLEREGI FMEDTACPYLABISMALGHLHQKGI I YRDLKPENIMLNHQGHVKLTD
                                                               GGELFTHLEREGIFLEDTACFYLAEITLALGHLHSQGIIYRDLKFENIMLSSQGHIKLTD
                                                                                                                               GKI FAMKVLKKAMI VRNAKDTAHTKABRNI LBEVKHPF I VDLI YAPQTGGKLYLI LBYLS
                                                                                                                                                                                                                GGVGPYELGMEHCÉKFÉISÉTŚVNRGPÉKÍRPEĆFELLRVLGKGGYGKVFQVRKVTGANT 116
                                                                                                                                                                                                                                                                                                MRRRRRRDGFYPAPDFRDREAEDMAGVFDIDLDQPEDAGSEDELE----EGGQLNESMDH 56
                                                                                                                                                                                                                                                                                                                                      MARGRRARGAGAA------MAAVFDLDLETEEGSEGEGEPELSPADACPLAE-LRA 49
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Matches

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Query Match Best Local Similarity

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US-08-749-902-7
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                                       Query Match
Best Local Similarity
Matches 351; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                  TELEPAX: 415-845-4166
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bandman, Olga
APPLICANT: Goli, Surya K.
APPLICANT: Hillman, Jennifer L.
TITLE OF INVENTION: NOVEL HUMAN SERINE/THREONINE
TITLE OF INVENTION: PROTEIN KINASES
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/749, FILING DATB: Filed Herewith PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                             TOPOLOGY: linear IMMEDIATE SOURCE: LIBRARY: GenBank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 3174 Port
CITY: Palo Alto
STATE: CA
COUNTRY: US
                                                                                                                                                                      LENGTH: 525 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94304
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   1 MARGRRARGAGAA-----
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3174 Porter Drive
                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IBM Compatible
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                                          66.8%; Score 1747; DB 2; 67.9%; Pred. No. 5.2e-137; Dive 49; Mismatches 79;
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-MAAVFDLDLETEEGSEGEGSPELSPADACPLAE-LRA 49
                                                                      Length 525;
                                             Indels
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                                            Gaps
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US-09-430-564-16
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APPLICANT:
APPLICANT:
                                                                                                                                                                      Match 66.8%;
Local Similarity 67.9%;
les 351; Conservative 49
 417
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                                                                         50 AGLEP----VGHYEEVELTETSVNVGPERIGPHCFELLRVLGKGGYGKVFQVRKVQGTNL 105
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                                                                                                                                1 MARGRRARGAGAA-------MAAVFDLDLETEEGSEGEGEBELSPADACPLAE-LRA
                                                        GGVGPYBLGMEHCEKFEISETSVNRGPEKIRPECFELLRVLGKGGYGKVFQVRKVTGANT
                                                                                                                 MRRRRRRDGFYPAPDFRDREAEDMAGVFDIDLDQPEDAGSEDELE----EGGQLNESMDH
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49;

Score 1747; DB 4; Length 525; Pred. No. 5.2e-137; 9; Mismatches 79; Indels 3

38;

Gaps

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116

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SOFTWARE: FastSEQ for W:
SEQ ID NO 16
LENGTH: 525
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16, Application US/09430564
Patent No. 6372467
                                                                                                                TITLE OF INVENTION: P5456K AND P8556K GENES, PROTEINS, TITLE OF INVENTION: PRIMERS, PROBES, AND DETECTION MET FILE REFERENCE: 00246/506002

CURRENT APPLICATION NUMBER: US/09/430,564

CURRENT FILING DATE: 1999-10-29

PRIOR APPLICATION NUMBER: 60/106,141

PRIOR PILING DATE: 1998-10-29

RIOR APPLICATION NUMBER: 60/106,141

PRIOR FILING DATE: 1998-10-29

NUMBER: 0580 ID NOS: 16
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                                                                                               FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                              Kay K. Lee-Fruman
Calvin J. Kuo
                                                                                                                                                                                                                                                                                                                                                              John Blenis
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APPLICANT: Stewart, Mary
APPLICANT: Kozma, Sarah
APPLICANT: Kozma, Sarah
APPLICANT: Thomas, George
TITLE OF INVENTION: Drosophila Melanogaster P7(
FILE REFERENCE: 4-20971/A
CURRENT APPLICATION NUMBER: US/09/817,310
CURRENT FILING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: 09/230,247
PRIOR FILING DATE: 1999-04-16
NUMBER OF SEQ ID NOS: 5
SOCTWARE: Patentin Ver. 2.0
SEQ ID NO 2
TENOMERIC 237
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US-09-817-310-2
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 637
TYPE: PRT
ORGANISM: Drosophila melanogaster
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279; Conserv
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p. 6534311
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                                                      GHNRAVDWWSLGALMYDMLTGSPPFTAENRKKTMDKIIRGKLALPPYLTPDARDLVKKFL
                                                                                                                                  LHSQGIIYRDLKPENIMLSSQGHIKLTDFGLCKESIHEGAVTHTFCGTIIYMAPBILVRS
                                                                                                                                                                           AVKHPFIVZAFQTDGKIXLILZYLSGGELFMHLEREGIFLEDTTCFYLSEIILALGH
                                                                                                                                                                                                           SVKHPFIVELAYAFQTGGKLYLILECLSGGELFTHLEREGIFLEDTACFYLABITLALGH
                                                                                                                                                                                                                                                 KDFELKKVLGKGGYGKVFQVRKTAGRDANKYFAMKVLKKASIVTNQKDTAHTRAERNILE
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 KRNPSQRIGGGPGDAADVQRHPFFRHMNWDDLLAWRVDPPFRPCLQSEEDVSQFDTRFTR 377
                                 GHGKAVDWWSLGALMFDMLTGVPPFTAENRKKTIETILKAKLNLPAYLTPEARDLVRRLM
                                                                                                       LHKLGIIYRDLKPENILLDAQGHVKLTDFGLCKEHIQEGIVTHTFCGTIEYMAPBILTRS
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                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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US-08-749-902-3

Sequence 3, Patent No.

Application US/08749902

GENERAL INFORMATION:

APPLICANT: Bandman, Olga
APPLICANT: Goli, Surya K.
APPLICANT: Hillman, Jennifer L.
TITLE OF INVENTION: NOVEL HUMAN SERINE/THREONINE
TITLE OF INVENTION: PROTEIN KINASES
NUMBER OF SEQUENCES: 8

CORRESPONDENCE ADDRESS

CITY: Palo Alto STATE: CA

STREET: ADDRESSEE:

3174 Porter Drive

INCYTE PHARMACEUTICALS,

RESULT 8

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US-09-430-564-12
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US-09-430-564-12
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SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 12, Appli
Patent No. 6372467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Kay K. Lee-Fruman
APPLICANT: Kay K. Lee-Fruman
APPLICANT: Calvin J. Kuo
TITLE OF INVENTION: PS456K AND P8556K GENES, PROTEINS,
TITLE OF INVENTION: PRIMERS, PROBES, AND DETECTION METHODS
FILE REFERENCE: 00246/506002
CURRENT APPLICATION NUMBER: US/09/430,564
CURRENT FILING DATE: 1999-10-29
PRIOR APPLICATION NUMBER: 05/106,141
PRIOR FILING DATE: 1998-10-29
PRIOR FILING DATE: 1998-10-29
PRIOR FILING DATE: 1998-10-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                               Local Similarity
241
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                                                             181
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                                                                                                                             LHSQGIIYRDLKPENIMLSSQGHIKLTDFGLCKESIHBGAVTHTFCGTIEYMAPEILVRS
                                                                                                                                                 LHSQGIIYRDLKPENIMLSSQCHIKLTDFGLCKESIHEGAVTHTFCGTIEYMAPEILVRS
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                                                                GHNRAVDWWSLGALMYDMLTGSPPFTAENRKKTMDKI IRGKLALPPYLTPDARDLVKKFL
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RESULT 9
US-09-430-564-6
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                                Sequence 6, Application Patent No. 6372467 GENERAL INFORMATION:
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Best Local Similarity
Matches 282; Conserv
   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 495 amino acids
TYPE: amino acid
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CLONE: Consens
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REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0150 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/7-FILING DATE: Filed Herewith PRIOR APPLICATION DATA:
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SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                  449
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                                                                                                                                                                                                                                                          GFTYVAPSVLDSIKEGFSFQPKLRSPRRLNSSPRVPVSPLKFSPFEGFRPSPSLPEPTEL
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                                                                                                                                                              PLPPLLPPPPPSTTAPLPIRPPSGTKKSKRGRGRPGR
                                                                                                                                                                                             PLPPLLPPPPPSTTAPLPIRPPSGTKKSKRGRGRPGR 495
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                                                                                                                                                                                                                                   GFTYVAPSVLDSIKEGFSFQPKLRSPRRLNSSPRVPVSPLKFSPFEGFRPSPSLPEPTEL
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John Blenis
Kay K. Lee-Fruman
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                                                                       US/09430564
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RESULT 10
US-09-031-295-2
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Best Local Simi
Matches 258;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application US/09031295
Patent No. 6326181
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
SEQ ID NO 6
                                                                                                                                                              ZIP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/031,295
FILING DATE: 26-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Calvin J. Kuo
TITLE OF INVENTION: P5956K AND P8586K GENES, PROTEINS,
TITLE OF INVENTION: PRINCES, PROBES, AND DETECTION METHODS
FILE REFERENCE: 00246/506002
CURRENT APPLICATION NUMBER: US/09/430,564
CURRENT FILING DATE: 1999-10-29
PRIOR APPLICATION NUMBER: 60/105,141
PRIOR PILING DATE: 1998-10-29
PRIOR FILING DATE: 1998-10-29
RIOR FILING DATE: 1998-10-29
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RIOR FILING DATE: 1998-10-29
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ORGANISM: Homo sapiens
                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMER: DE 197-08-173.8
FILING DATE: 28-FEB-1997
ATTORNEY/AGENT INFORMATION:
NAME: CRITICALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: LANG, Florian
APPLICANT: WALDEGGER, Tubingen
TITLE OF INVENTION: CELL VOLUME-REGULATED HUMAN KINASE H-SGK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 258
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ADDRESSEE: FOLEY & L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
NAME: Sandercock, Colin G. REGISTRATION NUMBER: 31,298 REFERENCE/DOCKET NUMBER: 05
                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     318 KRNPSQRIGGGPGDAADV 335
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      198 LHSQGIIYRDLKPENIMLSSQGHIKLTDFGLCKESIHEGAVTHTFCGTIEYMAPEILVRS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      138 SVKHPFIVBLAYAFQTGGKLYLILECLSGGBLFTHLBRBGIFLBDTACFYLABITLALGH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 SVKHPFIVELAYAPQTGGKLYLILECLSGGELFTHLEREGIFLEDTACFYLABITLALGH
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Similarity 100.0%;
58; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                 Washington
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                                                                                                                                                                                                                                                                                                                                                             U.S.A.
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                                                                                                                                                                                                                                  Release #1.0,
                                                                                                                                                                                         us/09/031,295
  058315/0123
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; Pred. No. 9.2e-105;
0; Mismatches 0;
                                                                                                                                                                                                                                      Version
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US-08-712-709-5
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 431 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 5, Application US/08712709
Patent No. 5863780
GENERAL INFORMATION:
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Best Local .
                                                  SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/712,
FILING DATE: Filed Herewith
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                    ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                              FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R
                                                                                                                                                 COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                     STREET: 3174 POI
CITY: Palo Alto
                                                                                                                                                                                                                                  COUNTRY: U.S.
ZIP: 94304
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
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                  Billings, Lucy
RATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KSPDSVLVTASVKEAAEAFLGFSYAPPT 427
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GSFGKVLLARHKABEV----FYAVKVLQKKAILKK-KEEKHIMSERNVLLKNVKHPFLVG 161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGYGKVFQVR-KVQGTNLGKIYAMKVLRKAKIVRNAKDTAHTRAERNI-LESVKHPFIVE 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EPELMMANPSP-----PPSPSQQ-----INLGPSS-NPHAKPSDFHFLKVIGK 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAKDDFMEIKSHVFFSLINWDDLINKKITPPFNPNVSGPNELRHFDPEFTEE-PVPNSIG
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                                                                                                                                                                                                                                                                                                              E: Incyte Pharmaceuticals,
3174 Porter Drive
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                                                                                                                                                                   3: Diskette
IBM Compatible
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44.3%; Pred. No. 5.7e-59;
tive 66; Mismatches 111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Janice
                                                                                            US/08/712,709
PF-0118
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US-09-111-444-5

Sequence 5, Application US/09111444
Patent No. 6045792

GENERAL INFORMATION:

APPLICANT: Au-Young, Janice
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: NOVEL HUMAN PROTEIN KINASES
NUMBER OF SEQUENCES: 9

CORRESPONDENCE ADDRESS

STREET:

Palo Alto

ς

ADDRESSEE:

E: Incyte Pharmaceuticals, 3174 Porter Drive

COUNTRY:

U.S.

94304

COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/111,
FILING DATE:

US/09/111,444

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette

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US-08-712-709-5
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
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400 KSPDSVLVTASVKEAAEAFLGFSYAPPT 427
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                                                                                                                                                                                                                                                                                                                                                                                                                  65 EPELMNANPSP------PPSPSQQ-----INLGPSS-NPHAKPSDFHFLKVIGK 106
                                 DSPDD---TALSESANQAFLGFTYVAPS 406
                                                                                                   GGPGDAADVQRHPFFRHMMDDLLAWRVDPPFRPCLQSBBDVSQFDTRFTRQTPV-----
                                                                                                                                      CLGAVLYEMLYGEPPFYSRNTAEMYDNILNKPLQLKPNITNSARHLLEGLLQKDRTKRL-
                                                                                                                                                                     SIGALMYDMITGSPPFTAENRKKTMDKIIRGKLALPPYLTPDARDLVKKPLKRNPSQRIG 326
                                                                                                                                                                                                          DIKPENILIDSQGHIVLTDFGLCKENIEHNSTTSTFCGTPEYLAPEVLHKQPYDRTVDWW
                                                                                                                                                                                                                                    DLKPENIMLSSQGHIKLTDFGLCKESIHBGAVTHTFCGTIBYMAPEILVRSGHNRAVDWW
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                                                                    GAKODFMEIKSHVFFSLINWDDLINKKITPPFNPNVSGPNDLRHFDPBFTEE-PVPNSIG
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Sequence 5, Application US/09541228
Patent No. 6232077
GENERAL INFORMATION:
APPLICANT: Au-Young, Janice
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: NOVEL HUMAN PROTEIN KI.
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       문
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US-09-541-228-5
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Best Local Similarity
Matches 172; Conserv
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INFORMATION FOR SEQ ID NO:
               COMPUTER
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LENGTH: 431 amino acids
TYPE: amino acid
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                              STREET: 3174 Port
CITY: Palo Alto
STATE: CA
COUNTRY: U.S.
ZIP: 94304
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLONE:
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             READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                        KSPDSVLVTASVKEAAEAFLGFSYAPPT 427
                                                                                                                                                                                                                                                                                                                                                                     DSPDD---TALSESANQAFIGETYVAPS 406
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                                                                                                                                                                                                                                                                                                                                                                                                      GAKDDFMEIKSHVFFSLINWDDLINKKITPPFNPNVSGPNDLRHFDPEFTEE-PVPNSIG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLGAVLYEMLYGLPPFYSRNTAEMYDNÍLNKPLQLKPNITNSARHLLEGLLQKDRTKRL-
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                                                                                                                                                                                                                                                                                                                                                                                                                                   GGPGDAADVQRHPFFRHMWWDDLLAWRVDPPFRPCLQSEEDVSQFDTRFTRQTPV-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Consensus
                                                                                           3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30.7%; Score 802.5; DB 3; 44.3%; Pred. No. 1.2e-58; ative 65; Mismatches 112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08/712,709
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                                                                                                                                                            KINASES
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                                                                                                                                                          KRESULT 14
US-08-712-709-9
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          Sequence 9, Application US/08712709
PATCHI NO. 5863780
GENERAL INFORMATION:
APPLICANT: Au Young, Janice
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: MOYEL, HUMAN PROTEIN KINASES
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
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US-09-541-228-5
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Best Local Similarity 44.3%;
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INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/541,228
FILING DATE:
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CLONE: Consensus
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TOPOLOGY: lir
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SOFTWARE: FASCSEQ Version 1.5
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COMPUTER: II
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                                                                                     382
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400 KSPDSVLVTASVKEAAEAFLGFSYAPPT 427
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             267 SLGALMYDMLTGSPPFTAENRKKTMDKIIRGKLALPPYLTPDARDLVKKFLKRNPSORIG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            222 DIKPENILLDSOGHIVLIDFGLCKENIEHNSTTSTFCGTPEYLAPEVLHKQPYDRTVDWW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      162 LHFSFQTADKLYFVLDYINGGELFYHLQRERCFLEPRARSYAAEIASALGYLHSLNIVYR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             147 LAYAFQTGGKLYLILECLSGGELFTHLEREGIFLEDTACFYLABITLALGHLHSQGIIYR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     107 GSFGKVLLARHKABEV----FYAVKVLQKKAILKK-KEEKHIMSERNVLLKNVKHPFLVG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                89 GGYGKVFQVR-KVQGTNLGKIYAMKVLRKAKIVRNAKDTAHTRAERNI-LESVKHPFIVB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65 EPELMNANPSP-----PPSPSQQ-----INLGPSS-NPHAKPSDPHFLKVIGK
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                                                                                     DSPDD---TALSESANQAFLGFTYVAPS 406
                                                                                                                                                                                    GAKDDFMEIKSHVFFSLINWDDLINKKITPPFNPNVSGPNDLRHFDPBFTEE-PVPNSIG 399
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; Pred. No. 1.2e-58; 
65; Mismatches 112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PF-0118
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Gaps

340 326 266 221

281

161 146 106 89

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MOLECULE TYPE: pe
; IMMEDIATE SOURCE;
; LIBRARY: GenBan
; CLONE: 294637
US-08-712-709-9
                                                                                       RESULT 15
US-09-111-444-9
                              Sequence 9, Application Patent No. 6045792 GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 430 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: 415-855-0555
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FILING DATE: Filed Herew.
ATTORNEY/AGENT INFORMATION:
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SOPTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0118 US
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                                                                                                                                                                                                                                                                                                         CLGAVLYEMLYGLPPFYSRNTAEMYDNILNKPLOL-KNITNSARHLLEGLLQKDRTKRL-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGYGKVFQVR-KVQGTNLGKIYAMKVLRKAKIVRNAKDTAHTRABRNI-LBSVKHPFIVB 146
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                                                                                                                                                                                               -SPDD---TALSESANQAFLGFTYVAPSVLDS 410
                                                                                                                                                                                                                                    GAKDDFWEIKSHIFFSLINWDDLINKKITPPFNENVSGPSDLRHFDPEFTEE-PVPSSIG
Au-Young, Janice
Guegler, Karl J.
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                                                                                                                                                              428
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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SEQUENCE CHARACTERISTICS:
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IMMEDIATE SOURCE:
LIBRARY: GenBar
CLONE: 294637
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TITLE OF INVENTION: NOVEL HUMAN PROTEIN KINASES
NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
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REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
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CITY: Palo Alto
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                                    RSPDSILVTASVKEAAEAFLGESYAPP--MDS
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3174 Porter Drive
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Search completed: June 30, 2004, 17:15:19 Job time : 25 secs

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Query Match 94.9%; Score 1724; D Best Local Similarity 100.0%; Pred. No. 0; Matches 1724; Conservative 0; Mismatches

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B9 CGACGGGCCCCCCGCGGCGCCCCCATGGCGGCCGTTTTTGGATTTTGGATTTTGGAGACC

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TOPOLOGY: linear
IMMEDIATE SOURCE:
LIBRARY: SXINBITO1
CLONE: 1869688
US-08-966-316-10
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Patent No. 5932445
GENERAL INFORMATION:
APPLICANT: Lal, Preeti
APPLICANT: Lal, Preeti
APPLICANT: Reddy, Roopa
APPLICANT: Matry, Lynn B.
APPLICANT: Matry, Lynn B.
APPLICANT: Matry, Preete
TITLE OF INVENTION: SIGNAL PEPTIDE - CONTAINING PROTEINS
NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
annowed: Theore a pharmaceuticals, Inc.
Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTESQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/966,316
FILING DATE: Herewith
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
FILING DATE:
FILING DATE:
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APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
FILING DATE
                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1637 base pairs
TYPE: nucleic acid
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3174 Porter Drive
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60.8%;
98.8%;
    Score 1104.4; DB 2;
Pred. No. 4.5e-249;
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Best Local Similarity 99.1%;
Matches 734; Conservative
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 1607 base pairs
TYPE: nucleic acid
STRANDENESS: single
TOPOLOGY: linear
INMEDIATE SOURCE:
CLONE: Consensus
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Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

COMPUTER: IBM Compatible

OPERATING SYSTEM: DOS

SOFTWARE: FastSEQ Version 2.0

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/749

FILING DATE: Filed Herewith

PRIOR APPLICATION NUMBER:

APPLICATION NUMBER:

APPLICATION NUMBER:

APPLICATION NUMBER:

APPLICATION NUMBER:

FILING DATE:

ATTORNEY/AGENT INFORMATION:

NAME: Billings, Lucy J.

REFERENCE/DOCKET NUMBER: PF-0:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
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APPLICANT: Bandman, Olga
APPLICANT: Goli, Surya K.
APPLICANT: Hillman, Jennifer L.
TITLE OF INVENTION: NOVEL HUMAN SERINE/THREONINE
TITLE OF INVENTION: PROTEIN KINASES
NUMBER OF SEQUENCES: 8
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ADDRESSEE: INCYTE PHARMACEUTICALS,
STREET: 3174 Porter Drive
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CITY: Palo Alto
STATE: CA
COUNTRY: US
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Pred. No. 3.7e-159;
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Patent No. 6500938
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FILING DATE: HEREWITH CLASSIFICATION:
PRIOR APPLICATION DATA: APPLICATION NUMBER: FILING DATE:
                                                                            ZIP: 94304
ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/016,434
                                                                                                                                                                                                                                                                                                                                              APPLICANT: Janice Au-Young
APPLICANT: Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION
TITLE OF INVENTION: PATHWAY GENE EXPRESSION
                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 1. CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                         ADDRESSEE: INCYTE PHARMACEUTICALS,
STREET: 3174 PORTER DRIVE
CITY: PALO ALTO
STATE: CALIFORNIA
                                                                                                                                                                                                           COUNTRY: USA
ZIP: 94304
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ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: 97,071
REFERENCE/DOCKET NUMBER: 985-055
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 845-4166
INFORMATION FOR SED ID NO: 126:
SEQUENCE CHARACTERISTICS:
LENGTH: 1607 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
LINEARY: COLNFETO2
LINEARY: COLNFETO2
CLONE: 1309709
US-09-016-434-126
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Best Local Similarity
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                                                        CAAGAGGGCCGTGGGCGTCCAGGGCG-TAGGAAGCCGGGTGGGGGGTAGCCCTT
                                                                            CAAGAGGGGCCGTGGGGGTCCAGGGCGCTAGGAAGCCGGGTGGGGGTGAGGGTAGCCCTT
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                                                                                                                                                                                                                                                                 CCTCAACAGTAGCCCCGGGGCCCGTCAGCCCCCCCAAGTTCTCCCCCTTTTGAGGGGTT
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Pred. No. 3.7e-159;
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; TOPOLOGY: linear
; INVEDIATE SOURCE;
; LIBRARY: BRAINOT
; CLONE: 1290913
US-09-016-434-118
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US-09-016-434-118
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Best Local S
Matches 646
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Patent No. 650093
                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 669 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA
TBLECOMYUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
TELEPAX: (650) 845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Janice Au-Young
APPLICANT: Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF SIGNALING
TITLE OF INVENTION: PATHMAY GENE EXPRESSION
NUMBER OF SEQUENCES: 1490
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: HEREWITH
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
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STATE: CALIFORNIA
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646; Conserv
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                                         CAGCGTGAACGTTGGCCCAGAGCGCATCGGGCCCCACTGCTTTGAGCTGCTGCGTGTGCT
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                                                                 CAGCGTGAACGTTGGCCCAGAGCGCATCGGGCCCACTGCTTTGAGCTGCTGCGTGTGCT
                                                                                             GTTGAGGGCAGCTGGCCTAGAGCCTGTGGGACACTATGAAGAGG-----GACTGAGAC
                                                                                                                     AGGCAGCGAGGGGAGGCGAGCCAGAGCTCAACNCCGCGGACGCATGTNCCCTTGCCGA
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95.8%;
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Pred. No. 1e-129;
0; Mismatches 1:
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                                                                                                                                                                                                                                                       18;
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; ORGANISM: Drosophila melanogaster
US-09-817-310-1
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US-09-817-310-1
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Best Local Similarity
Matches 718; Conserv
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APPLICANT: Stewart, Mary
APPLICANT: Kozma, Sarah
APPLICANT: Thomas, George
ITILE OF INVENTION: Drosophila Melanogaster P70 S6
ITILE OF INVENTION: Drosophila Melanogaster P70 S6
ITILE REFERENCE: 4-20971/A
CURRENT APPLICATION NUMBER: US/09/817,310
CURRENT FILING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: 09/230,247
PRIOR PILING DATE: 1999-04-16
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
LENGTH: 2556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/09817310 Patent No. 6534311 GENERAL INFORMATION:
                                                                                                                                                                                                        321 TGCTGCGTGTGCTGGGCAAGGGGGCTATGGCAAGGTGTTCCAGGTGCGAAAGGTGCAAG
                                                                                                                                                                                                                                                         649 AGCTCTGCGAGGAGAATGTTAATCCAGGTAAAATCAAGCTGGGACCCAAGGACTTTGAGC
                                                                                                                                                                                                                                                                                                          261 AGCTGACTGAGACCAGCGTGAACGTTGGCCCAGAGCGCATCGGGCCCCACTGCTTTGAGC
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ATGCCAAGGACACACACACACGGGCTGAGCGGAACATTCTAGAGTCAGTGAAGCACC
                                                   GACGAGATGCTAACAAATATTTTGCCATGAAGGTGCTCAAAAAGGCATCCATTGTGACCA
                                                                                         TCANGANGGTCCTTGGCANAGGCGGTTATGGCANAGTATTTCAGGTGCGCAAGACCGCTG
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Pred. No. 6.6e-111
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APPLICANT: Ole THASTRUP, et al.
TITIE OF INVENTION: A Method for Extracting Quartifle OF INVENTION: On A Cellular Response FILE REFERENCE: 3759-0110P
CURRENT APPLICATION NUMBER: US/09/417,197
CURRENT PILING DATE: 1999-10-07
NUMBER OF SEQ ID NOS: 143
SOPTWARE: PatentIn version 3.0
SEQ ID NO 70
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US-09-417-197-138
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Patent No. 6518021
GENERAL INFORMATION:
APPLICANT: Ole THASTRUP, et al.
TITLE OF INVENTION: A Method For Extracting Quantitative Information |
TITLE OF INVENTION: On A Cellular Response
FILE REFERENCE: 3759-0110P
CURRENT APPLICATION NUMBER: US/09/417,197
CURRENT FILING DATE: 1999-10-07
NUMBER OF SECULATION SECULATION OF SECULATION NUMBER: US/09/417,197
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SEQ ID NO 138
LENGTH: 2184
TYPB: DNA
ORGANISM: Artificial Sequence
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Pred. No. 1e-51;
0; Mismatches 384;
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; LENGTH: 2610
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (199)..(1641)
US-09-212-771-1
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US-09-212-771-1
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APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0034
CURRENT APPLICATION UNMERR: US/09/212,771
CURRENT FILING DATE: 1998-12-16
NUMBER OF SEQ ID NOS: 47
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Best Local Similarity
Matches 517; Conserv
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Patent No. 5958773
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Pred. No. 1.1e-51;
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TITLE OF INVENTION: Screening Method
FILE REFERENCE: 4-2068)A9.70684/PCT
CURRENT APPLICATION UNUMER: US/09/091,058
CURRENT FILING DATE: 1998-06-10
EARLIER APPLICATION UNMER: PCT/SE96/04814
EARLIER FILING DATE: 1996-11-05
EARLIER FILING DATE: 1995-12-15
EARLIER FILING DATE: 1995-12-15
EARLIER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.0
LENGTH: 2610
TYPE: DNA
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; LOCATION: (199)..(1641)
US-09-091-058-1
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US-09-091-058-1
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Patent No. 6054285
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                                                                                                                                                                                                                                                                                   ORGANISM: Homo
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CANT: Cocks, Benjamin G.
CANT: Susan G. Stuart
CCANT: Jeffrey J. Seilhamer
COF INVENTION: COMPOSITION
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LIBRARY: GENBANK
CLONE: g190827
US-09-023-655-1206
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 517; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPAX: (650) 845-410
INFORMATION FOR SEQ ID NO:
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STREET: 31...
CITY: PALO ALTO
CTATE: CALIFORNIA
TTSA
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LENGTH: 2610 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAMS: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: HEREWITH
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Word Perfect 6.1 for Windows/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME:
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                                          CAAGCTGGAGAACCTCATGCTGGACAAGGACGGGCACATTAAGATCACAGACTTCGGGCT
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                                                                     CAAGCCCGAGAACATCATGCTCAGCAGCCAGGCCACATCAAACTGACCGACTTTGGACT
                                                                                                  TGAGATTGTGTCAGCCCTGGACTACCTGCACTCGGAGAAGAACGTGGTGCTACCGGGACCT
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Pred. No. 1.1e-51;
0; Mismatches 384;
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APPLICANT: Brett P. Monia
APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0035
CURRENT APPLICATION UNMERS: US/09/256,465
CURRENT FILING DATE: 1999-02-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 47 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/09256465 Patent No. 6043090
                                                                                                                                                                                                                                                                                                                               Matches 569;
                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: CDS
LOCATION: (88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Homo sapiens FEATURE:
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                                          CATTGCCAAGG---ATGAAGTCGCTCACACAGTCACCGAGAGCCCGGGTCCTCCAGAACAC
                                                                                                                                                                                   G-----AAGGCCACTGGCCGCTACTACGCCATGAAGATCCTGCGAAAGGAAGTCAT
                                                                                                                                                                                                                                CTTCGACTATCTCAAACTCCTTGGCAAGGGAACCTTTGGCAAAGTCATCCTGGTGCGGGA
                                                                                       TGTGCGCAATGCCAAGGACACACACACACGGGCTGAGCGGAACATTCTAGAGTCAGT
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Pred. No. 4.1e-50;
0; Mismatches 472;
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RESULT 13
US-09-167-322-3
US-09-167-322-3
; Sequence 3, Application US/09167322
; Patent No. 6365151
; GENERAL INFORMATION:
; GENERAL INFORMATION:
APPLICANT: Allegheny University of the Health
Sciences, Halpern, Michael S.
England, James M.
TITLE OF INVENTION: CANCER VACCINE
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Seidel, Gonda, Lavorgna & Monaco, P.C.
STREET: Suite 1800, Two Penn Center Plaza
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FILING DATE: «UDKNOWN»
ATTORNEY/AGENT INFORMATION:
NAME: MODACO, Daniel A.
REGISTRATION NUMBER: 30,480
REFERENCE/DOCKET NUMBER: 7933-33 PC
TELEOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-8383
TELEPHANE: (215) 568-8383
TELEPHANE: (215) 568-8383
INFORMATION POR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 14.0%;
Best Local Similarity 54.0%;
                                                1008
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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                                                GAAAACCTTCTGTGGGACCCCGGAGTACCTGGCGCCTGAGGTGCTGGAGGACAATGACTA
                                                                                                                                                              CCACATCAAACTGACCGACTTTGGACTCTGCAAGGAGTCTATCCATGAGGGCGCCGTCAC
      CAACCGGGCTGTGGACTGGTGGAGCCTGGGGGCCCTGATGTACGACATGCTCACTGGATC
                                                                                        TCACACCTTCTGCGGCACCATTGAGTACATGGCCCCTGAGATTCTGGTGCGCAGTGGCCA
                                                                                                                                      CCACATCAAGATCACTGACTTTGGCCTCTGCAAAGAGGGCATCAGTGACGGGGCCACCAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 1599 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Philadelphia
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Pred. No. 4.1e-50;
0; Mismatches 472;
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RESULT 14
US-09-023-655-1004
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ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-C
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cocks, Benjamin G.
APPLICANT: Susan G. Stuart
APPLICANT: Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 1508
                                                                                                                                                                                                                                   SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023,655
                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FLOPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                               PRIOR APPLICATION DATA:
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                                                                                                                     CLASSIFICATION:
                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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T: 3174 PORTER DRIVE
PALO ALTO
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; INFORMATION FOR SEQ ID NO: 10
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1599 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
; IMMEDIATE SOURCE:
; LIBRARY: GENBANK
; CLONE: 9178325
US-09-023-655-1004
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                                     GTTCTTCCTCAGCATCAACTGGCAGGACGTGGTCCAGAAGAAGCTCCTGCCACCCTTCAA 1367
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Pred. No. 4.1e-50;
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Search completed: July 3, 2004, 10:10:31 Job time: 153 secs

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Result
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MODEL-frame+ p2n.model -DEV=xlh
-Q-/Cgn2 _1/USPTO_Spoot/US09762258/runat_30062004_165608_13618/app_query.fasta_1.647
-DB=Issued_Patents_NA -QPMT-fastap_-SUFFIX=rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS-bits -START=1 -END=-1 -MATRIX=biosum62 -TRANS-buman40.cdi
-LIST=45 -DOCALICN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM=ext -HBAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=US09762258_@CGN 1 1 69 @runat_30062004_165608_13618 -NCFU-6 -ICPU-3
-NO_WMAP_LARGEQUERY -NEG_SCORES-0 -MAIT_-DSPBJOCKS=100 -LOOGLOG
-NO_WMAP_LARGEQUERY -NGS_SCORES-0 -MAIT_-DSPBJOCK=100 -LOOGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOD=10 -YGAPEXT=0.5 -DELOP=6 -DEPLEXT=7
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13	e 1191, A	e 1, Appl	e 3, Appl	s 52, App	1362, A	e 1. Appl	e 14, App	e 35, App	o. 526646	e 5, Appl	e 3, Appl	e 1361, F	e 3, Appl	e 1471, ;	e 1433, F	e 72, App	e 48, App	e 1, App]	e 1, Appl	e 24, App	e 25,	e 25, Api	e 25, Apr	e 1206, <i>i</i>	e 1, Appl	e 1, App]	e 138, Ar	e 70, Apr	e 24, App	e 25, Apı	Sequence 1004, Ap	e s

## ALIGNMENTS

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                                                US-09-762-258-2 (1-495) x US-09-430-564-1 (1-1732)
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US-09-430-564-1
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SEQ ID NO 1
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Patent No. 6372467
GENERAL INFORMATION:
APPLICANT: John Blenis
APPLICANT: Kay K. Lee-Fruman
APPLICANT: Calvin J. Kuo
                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: P5485K AND P8586K GENES, PROTEINS, ITTLE OF INVENTION: PRIMERS, PROBES, AND DETECTION MET PILE REFERENCE: 00246/506002

CURRENT APPLICATION NUMBER: US/09/430,564

CURRENT FILING DATE: 1999-10-29

PRIOR APPLICATION NUMBER: 60/106,141

PRIOR FILING DATE: 1998-10-29

RUMBER OF SEQ ID NOS: 16
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ORGANISM: Homo:
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FOR	ATTERNATION OF THE PROPERTY OF
	325 IleGlyGlyProGlyAspAlaAlaAspValGlmArgHisProPhePheArgHisMet 344
REGISTRATION REFERENCE/DO TELECOMMUNICAT	305 LeuThrProAspAlaArgAspLeuValLysLysPheLeuLysArgAsnProSerGlnArg 324
; APPLICATION; FILING DATE; FILING DATE; ATTORNEY/AGENT; NAME; Billi;	285 GluAsnArgLysLysThrMetAspLysIleIleArgGlyLysLeuAlaLeuProProTyr 304 
; APPLICATION ; FILING DATE; ; CLASSIFICATION ; PRIOR APPLICATION	265 TrpTrpSerLeuGlyAlaLeuMetTyrAspMetLeuThrGlySerProProPheThrAla 284
COMPUTER: I OPERATING SY SOFTWARE: F CURRENT APPLIC	245 ThrileGluTyrMetAlaProGluIleLeuValArgSerGlyHisAsnArgAlaValAsp 264 
COUNTRY: US ZIP: 94304 ; COMPUTER READA ; MEDIUM TYPE:	225 AspPheGlyLeuCysLysGluSerIleHisGluGlyAlaValThrHisThrPheCysGly 244
; ADDRESSEE: STREET: 317; CITY: Palo.; STATE: CA	205 TyrArgAspLeuLysProGluAsnIleMetLeuSerSerGlnGlyHisIleLysLeuThr 224
9 S S	185 CysPheTyrLeuAlaGluIleThrLeuAlaLeuGlyHisLeuHisSerGlnGlyIleIle 204
APPLICANT: LA APPLICANT: Au APPLICANT: Au APPLICANT: Re APPLICANT: Mi	165 SerGlyGlyGluLeuPheThrHisLeuGluArgGluGlyIlePheLeuGluAspThrAla 184 
US-08-966-316-10 US-08-966-316-10 ; Sequence 10, Appl ; Patent No. 593244 ; GENERAL INFORMAT	145 ValGluteuAlaTyrAlaPheGlnThrGlyGlyLysLeuTyrLeuIleLeuGluCysLeu 164 
μ.	125 ASpThrAlaHisThrArgAlaGluArgAsnIleLeuGluSerValLysHisProPheIle 144
1389	105 LeuGlyLysIleTyrAlaMetLysValLeuArgLysAlaLysIleValArgAsnAlaLys 124 
1329 (	85 ValleuGlyLysGlyGlyTyrGlyLysValPheGlnValArgLysValGlnGlyThrAsn 104 
Db 1269 AGGCG	65 GIUTRESETVALASMVAIGLYPEOGLUARGILGGLYPEOHISCYSPHEGIULGULGUAGG 84 [
Qy 425 ArgAr	GCCGAGTTGAGGCCTGGCCTAGAGCCTGGGACACTATGAAGAGGTGGAGCTGACT
	5 AlaGluLeuArgAlaAlaGlyLeuGluProValGlyHisTyrGluGluValGluLeuThr
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Au-Young, Janice
Reddy, Roopa
Murry, Lynn E.
Mathur, Preete
WENTION: SIGNAL PEPTIDE - CC
EQUENCES: 18
NCE ADDRESS:
INCE ADDRESS:
INCE ADDRESS:
INCYTE Pharmaceuticals, II
3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                             ENT INFORMATION:
111.ngs, Lucy J.
10N NUMBER: 96,749
/DOCKET NUMBER: PF-0424 US
CATION INFORMATION:
650-855-0555
650-845-4166
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pAlaArgAspLeuValLysEheLeuLysArgAsnProSerGinArgIleGlyGlyGl
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Patent No. 6534311

GENERAL INFORMATION:

APPLICANT: Kozmat, Mary

APPLICANT: Kozmat, Sarah

APPLICANT: Kozmat, Sarah

APPLICANT: Thomas, George

FITTHE OF INVENTION: Drosophila Melanogaster P70 S6 Kinase

FILE REFERENCE: 4-20971/A

CURRENT APPLICATION NUMBER: US/09/817,310

CURRENT FILING DATE: 2001-03-26

PRIOR APPLICATION NUMBER: 09/230,247

PRIOR FILING DATE: 1999-04-16

NUMBER OF SEQ ID NOS: 5

CONTROLL SECTION OF SE
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                                              AlaCysProLeuAlaGluLeuArgAlaAlaGlyLeuGluProValGlyHisTyrGluGlu
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Matches:
Conservative;
Mismatches:
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361 CysLeuGlnSerGluGluAspValSerGlnPheAspThrArgPheThrArgGlnThrPro 380	321 ProSerGlnArgIleGlyGlyGlyProGlyAspAlaAlaAspValGlnArgHisProPhe 340	ProPheThrAlaGluAsnArgLysLysThrMetAspLysIleIleArgGlyLysLeuAla	241 ThrPheCysGlyThrIleGluTyrMetAlaProGluIleLeuValArgSerGlyHisAsn 260	201 GlnGlyIleIleTyrArgAspLeuLysProGluAsnIleMetLeuSerSerGlnGlyHis 220	161 LeuGluCysLeuSerGlyGlyGluLeuPheThrHisLeuGluArgGluGlyIlePheLeu 180.	121 ArgAsmālalysāspīhrālaHisThrārgālaGluārgāsmīleLeuGluSerVallys 140	61 ValGluLeuThrGluThrSerValAsnValGlyProGluArglleGlyProHisCysPhe 80
				ADDRESS OF THE STREET			
TELECOMMUNICATION INFORMATION: TELEPHONE: 415-855-0555 TELEPAX: 415-845-4166 INFORMATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS: SEQUENCE CHARACTERISTICS: TYPE: nucleic acid STRANDEDNESS: single TYPE: nicel single TOPOLOGY: linear IMMEDIATE SOURCE: CLONE: Consensus US-08-749-902-4	APPLICATION NUMBER: US/08/149,302 FILING DATE: Filed Herewith PRIOR APPLICATION DATA: APPLICATION NUMBER: INTORNEY/AGENT INFORMATION: NAME: Billings, Lucy J. REGISTRATION NUMBER: 36,749 REFERENCE/DOCKET NUMBER: PF-0150 US	COUNTRY: US ZIP: 94304 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette COMPUTER: IBM Compatible COMPUTER: PastSEQ Version 2.0 CURRENT APPLICATION DATA:	: TITLE OF INVENTION: MOYEL HUMAN SERING/IRROWNINE : TITLE OF INVENTION: PROTEIN KINASES : NUMBER OF SEQUENCES: 8 : CORRESPONDENCE ADDRESS: 8 : ADDRESSEE: INCYTE PHARMACEUTICALS, INC. : STREET: 3174 Porter Drive : CITY: Palo Alto : STREET: OF Alto	902-4 4, Application 5, 5985635 5. SPROGRATION: CANT: Bandman, CANT: Goli, Sur CANT: Hillman,		449 PROSETLEUPROGLUPROTRINGLU	421 LeuargSerProArgArg-LeuAsmSerSerProArgValProValSer

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Best Local Similarity:
Query Match:
DB:
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nArgHisProPhePheArgHisMetAsnTrpAspAspLeuLeuAlaTrpArgValAspPr
                                                                            eLeuLysArgAsnProSerGlnArgIleGlyGlyGlyProGlyAspAlaAlaAspValGl
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                                                                                                                                     GGGCAAGCTGGCACTNCCCCCCTACCTCACCCCAGATGCCCGGGACCTTGTCAAAAAGTT
                                                                                                                                                                                                           CACTGGATCGCCGCCTTTACCGCAGAGAACCGGAAGAAACCATGGATAAGATCATCAG
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Patent No. 6500938

APPLICANT: Janice Au-Young
APPLICANT: Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF
TITLE OF INVENTION: PATHWAY GENE EXPRESSION
NUMBER OF SEQUENCES: 1490
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHADMAGENTE.
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US-09-016-434-126
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1607 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                           CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650, 855-0555
                                                                                                                                                                                                                                                                                                               SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/016,434
FILING DATE: HEREWITH
CLASSIFICATION:
                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                               TELEPAX:
                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Best Local Similarity:
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; LIBRARY: COLMFET02
; CLONE: 1309709
US-09-016-434-126
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NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0
TELECONMUNICATION INFORMATION:
TELEPRIONE: (650) 855-0555
TELEPAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: 118:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 118, Application US/09016434
Patent No. 6500938
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                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: Word Perfect 6.1 for Windows/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION UNBEER: US/09/016,434
                                                                                                                                                                                FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Janice Au-Young
APPLICANT: Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION
TITLE OF INVENTION: PATHWAY GENE EXPRESSION
NUMBER OF SEQUENCES: 1490
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
TOTAL OF THE PROPERTY OF TH
                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 3174 PORTER DRIVE
CITY: PALO ALTO
STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                       FILING DATE:
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Best Local Similarity:
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       Sequence 1, Application US/09031295
Patent No. 6326181
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IMMEDIATE SOURCE:
LIBRARY: BRAIN
CLONE: 1290913
APPLICANT:
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STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                              ValGluLeuAlaTyrAlaPheGlnThrGlyGlyLysLeuTyrLeuIleLeuGluCysLeu
                                                                                                                                  AspPheGlyLeuCysLys
                                                                                                                                                                                  TyrArgAspLeuLysProGluAsnIleMetLeuSerSerGlnGlyHisIleLysLeuThr
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LANG, Florian
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90.71%
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36.04%
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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Best Local Similarity:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Versi
SOFTWARE: Patentin Release #1.0, Versi
APPLICATION NUMBER: US/09/031,295
FILING DATE: 26-FEB-1998
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 197-08-173.8
FILING DATE: 28-FEB-1997
AFTORNEY/AGENT INFORMATION:
NAME: Sandercock, Colin G.
REGISTRATION NUMBER: 058315/0123
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (202) 672-5399
INFORMATION FOR SEQ ID NO:
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FEATURE:
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TITLE OF INVENTION: CELL VOLUMI
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
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CITY: Washington
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                                                                nucleic acid
HisThrArgAlaGluArgAsnIle---LeuGluSerValLysHisProPheIleValGlu 146
                                                                                                                                                                    TOGTCC---AATCCTCATGCTAAACCATCTGACTTTCACTTCTTGAAAGTGATCGGAAAG
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                                                TTCTATGCAGTCAAAGTTTTACAGAAGAAAGCAATCCTGAAAAAG---AAAGAGGAGAAG
                                                                                                                                     GlyGlyTyrGlyLysValPheGlnValArg---LysValGlnGlyThrAsnLeuGlyLys
                                                                                                                                                                                                      GluArgIleGlyProHisCys-----
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Matches:
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RESULT 8
US-08-712-709-6
                                                                                                                                                                                           Sequence 6, Application US/08712709 Patent No. 5863780
                                                                                                    GENERAL INFORMATION:
APPLICANT: Au-Young, Janice
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: NOVEL HUMAN PROTEIN KINASES
                                                                      CORRESPONDENCE ADDRESS:
                                                                                         NUMBER OF SEQUENCES:
   STREET:
CITY: F
STATE:
                                                     ADDRESSEE:
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                                 E: Incyte Pharmaceuticals,
3174 Porter Drive
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DB:
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Best Local Similarity:
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REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEPAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/712,
FILING DATE: Filed Herewith
ATTORNEY/AGENT INFORMATION:
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ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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IMMEDIATE SOURCE:
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STRANDEDNESS: single
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                                                                                                                                                                                           HisThrArgAlaGluArgAsmIle---LeuGluSerValLysHisProPheIleValGlu 146
                                                                                                                                                                                                                                                                TTCTATGCAGTCAAAGTTTTACAGAAGAAAGCAATCCTGAAAAAG---AAAGAGGAGAAG 432
TyrLeuAlaGluIleThrLeuAlaLeuGlyHisLeuHisSerGlnGlyIleIleTyrArg
                                                    LeuAlaTyrAlaPheGlnThrGlyGlyLysLeuTyrLeuIleLeuGluCysLeuSerGly 166
                                                                                                                                                                                                                                                                                          IleTyrAlaMetLysValLeuArgLysAlaLysIleValArgAsnAlaLysAspThrAla 127
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Matches:
Conservative:
Mismatches:
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RESULT 9
US-09-111-444-6
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Patent No. 6045792
GENERAL INFORMATI
                                                                                               ZIP: 94304
COMPUTER READABLE FORM:
COMPUTER: Diskette
COMPUTER: IBM Compatibl
OPERATING SYSTEM: DOS
                                  SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/111,444
FILING DATE:
 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Ph
                                                                                                                                                                                                                                                                                                                            ENERAL INFORMATION:
APPLICANT: Au-Young, Janice
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: NOVEL HUMAN
                                                                                                                                                                                                                                   STREET: 3174 Por
CITY: Palo Alto
                                                                                                                                                                                                 COUNTRY:
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                                                                                                                                                                                                                                                        3174 Porter Drive
                                                                                                                                                                                                   U.S.
                                                                                                         IBM Compatible
SYSTEM: DOS
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 08/712,709
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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ATTORNEY/AGENT INFORMATION:
ANAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415.855-0555
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                      GlyLeuCysLysGluSerIleHisGluGlyAlaValThrHisThrPheCysGlyThrIle
GGACTCTGCAAGGAGAACATTGAACACAACAGCACAACATCCACCTTCTGTGGCACGCCG
                                                                                           AspLeuLysProGluAsnIleMetLeuSerSerGlnGlyHisIleLysLeuThrAspPhe
                                                                                                                                             TATGCTGCTGAAATAGCCAGTGCCTTGGGCTACCTGCATTCACTGAACATCGTTTATAGA
                                                                                                                                                                    TyrLeuAlaGluIleThrLeuAlaLeuGlyHisLeuHisSerGlnGlyIleIleTyrArg
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PATENT NO. 6232077
GENERAL INFORMATION:
APPLICANT: Au-Young, Janice
APPLICANT: Guegler, Karl J.
APPLICANT: Hawkins, Phillip R.
TITLE OF INVENTION: NOVEL HUMAN PROTEIN KINASES
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
INFORMATION FOR SEQ ID NO:
                                  PILING DATE:
ATTORNEY AGENT INFORMATION:
ATTORNEY AGENT INFORMATION:
NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
THE FERENCE AGENT AGENT
                                                                                                                                 FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                              SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/541
                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
OPERATING SYSTEM: DO
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                                                                                                                                                                                                                                                                                         COUNTRY: U.S. ZIP: 94304
                           TELEPHONE: 415-855-0555
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                TGCCIGGGAGCIGICITGIAIGAGARGCIGIATGGCCIGCGCCITITIAIAGCCGAAAC
                                          SerLeuGlyAlaLeuMetTyrAspMetLeuThrGlySerProProPheThrAlaGluAsn
                                                                                             GluTyrMetAlaProGluIleLeuValArgSerGlyHisAsnArgAlaValAspTrpTrp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                           TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Janice Au-Young
APPLICANT: Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF SIGNALING
TITLE OF INVENTION: PATHWAY GENE EXPRESSION
NUMBER OF SEQUENCES: 1490
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,
              IMMEDIATE SOURCE:
                                                                                                                                             REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0002
TELECOMMUNICATION: INFORMATION:
TELEPHONE: (650) 855-0555
                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS CURRENT APPLICATION DATA:
LIBRARY:
                                          LENGTH: 2311 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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CLASSIFICATI
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CITY: PALO ALTO
                               TOPOLOGY:
                                                                             LENGTH:
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US-09-016-434-772
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                 AATTCCGCAAGACACCTCCTGGAGGGCCTCCTGCAGAAGGACAGGACAAAGCGGCTC---
                                                                                                                                                TECCTEGGAGETETTTATGAGATGETGTATGGECTTTTTATAGECGAAAC
                                             ProAspAlaArgAspLeuValLysLysPheLeuLysArgAsnProSerGlnArgIleGly 326
                                                                               ACAGCTGAAATGTACGACAACATTCTGAACAAGCCTCTCCAGCTGAAACCAAATATTACA
                                                                                                                                                                               SerLeuGlyAlaLeuMetTyrAspMetLeuThrGlySerProProPheThrAlaGluAsn
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Sequence 1, Application US/09256465
PAtent No. 6043090
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert
ITILE OF INVENTION: ANTISENSE MODULATION OF AKT
FILE REFERENCE: RTS-0035
CURRENT APPLICATION NUMBER: US/09/256,465
CURRENT FILING DATE: 1999-02-23
NUMBER OF SEQ ID NOS: 47
SEQ ID NO 1
SEQ ID NO 1
LENGTH: 1599
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Best Local Similarity:
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                                                               ArgAsnAlaLysAsp---ThrAlaHisThrArgAlaGluArgAsnIleLeuGluSerVal
                                                                                                              ----AAGGCCACTGGCCGCTACTACGCCATGAAGATCCTGCGAAAGGAAGTCATCATT
                                                                                                                                         GlnGlyThrAsnLeuGlyLysIleTyrAlaMetLysValLeuArgLysAlaLysIleVal 120
                                                                                                                                                                            GACTATCTCAAACTCCTTGGCAAGGGAACCTTTGGCAAAGTCATCCTGGTGCGGGAG---
                                                                                                                                                                                                GluLeuLeuArgValLeuGlyLysGlyGlyTyrGlyLysValPheGlnValArgLysVal
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                                                                                                                                                          Sequence 3, Application US/09167322
Patent No. 6365151
GENERAL INFORMATION:
                                                                           APPLICANT: Allegheny University of the Sciences, Halpern, Michael S England, James M.
TITLE OF INVENTION: CANCER VACCINE NUMBER OF SEQUENCES: 14
 CORRESPONDENCE ADDRESS:
ADDRESSE: Seidel, Gonda, Lavorgna & Monaco,
STREET: Suite 1800, Two Penn Center Plaza
CITY: Philadelphia
STATE: PA
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Best Local Similarity:
Query Match:
DB:
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SEQUENCE CHARACTERISTICS:
LENGTH: 1599 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
STRANDENNESS: single
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PRIOR APPLICATION DATA;
APPLICATION UNMBER: PCT/US97/00582
PILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 19102
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-POS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE DESCRIPTION: SEQ
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REGISTRATION NUMBER: 30,480
REFERENCE/DOCKET NUMBER: 79:
TELECOMMUNICATION INFORMATION:
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  SerGlnGlyIleIleTyrArgAspLeuLysProGluAsnIleMetLeuSerSerGlnGly
                                        ACAGAGGAGCGGGCCCGGTTTTATGGTGCAGAGATTGTCTCGGCTCTTGAGTACTTGCAC
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TELEPAX: (215) 568-5549
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                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1004, Application US/09023655
Patent No. 6607879
GENERAL INFORMATION:
            COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
COMPUTER: PM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTYWARE: Word Perfect 6.1 for Wind
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023,655
                                                                                                                                                                                                                                                                                        APPLICANT: Cocks, Benjamin G. APPLICANT: Susan G. Stuart APPLICANT: Jeffrey J. Seilhamer TITLE OF INVENTION: COMPOSITION FOR TITLE OF INVENTION: EXPRESSION NUMBER OF SEQUENCES: 1508
                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PR
                                                                                                                                                                           CITY: PASTATE: COUNTRY:
                                                                                                                                                            ZIP:
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DATE:
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LIBRARY: GENEAL
CLONE: g178325
US-09-023-655-1004
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NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0/
TELECOMMUNICATION INFORMATION:
TELECHONE: (650) 85-0555
TELEFAX: (650) 85-4166
INFORMATION FOR SEQ ID NO: 1004:
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                     HisIleLysLeuThrAspPheGlyLeuCysLysGluSerIleHisGluGlyAlaValThr
                                                                                                                                 ACAGAGGAGCGGGCCCGGTTTTATGGTGCAGAGATTGTCTCGGCTCTTGAGTACTTGCAC
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                                                                                               SerGlmGlyIleIleTyrArgAspLeuLysProGluAsnIleMetLeuSerSerGlmGly
 CACATCAAGATCACTGACTTTGGCCTCTGCAAAGAGGGCATCAGTGACGGGGCCACCATG 1008
                                                                   TCGCGGGACGTGGTATACCGCGACATCAAGCTGGAAAACCTCATGCTGGACAAAGATGGC
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Matches:
Conservative:
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; Sequence 25, Application US/076883520; Patent No. 5527896; GENERAL INFORMATION:
APPLICANT: Wigler, Michael H.
APPLICANT: Wigler, Michael H.
APPLICANT: Colicelli, John J.
TITLE OF INVENTION: Cloning by Co.
TITLE OF INVENTION: Processes
NUMBER OF SEQUENCES: 57
CORRESPONDENCE ADDRESS:
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US-07-688-352C-25
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APPLICATION NUMBER: US/07/
FILING DATE: 19310419
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
FILING DATE: 20-APR-1990
ATTORNEY/AGENT INFORMATION:
NAME: BOTUN, Michael F.
                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM CO compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                 STREET: Two First National Plaza,
STREET: Street
CITY: Chicago
STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Marshall, O'Toole, Gerstein, Murray ADDRESSEE: Bicknell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GACCCCAAGCAGAGGCTTGGTGGGGGGCCCAGCGATGCCAAGGAGGTCATGGAGCACAGG 1308
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTGCCCTTCTACAACCAGGACCACGAGCGCCTCTTCGAGCTCATCCTCATGGAAGAGATC 1188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ProProPheThrAlaGluAsnArgLysLysThrMetAspLysIleIleArgGlyLysLeu 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGCCGGCCGTGGACTGGTGGGGGGCTGGGTCATGTACGAGATGATGTGCGGCCGC 1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACCCACTICCCCCAGTICICCTACTCGGCCAGCATCCGCGAGTGAGCAGTCTGCCCACGC 1548
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ProCysLeuGlnSerGluGluAspValSerGlnPheAspThrArgPheThrArgGln---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTCTTCCTCAGCATCAACTGGCAGGACGTCGTCCAGAAGAAGCTCCTGCCACCCTTCAAA 1368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AlaLeuProProTyrLeuThrProAspAlaArgAspLeuValLysLysPheLeuLysArg
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DB:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 25447
REFERENCE/DOCKET NUMBER: 2780
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 346-5750
TELEPHONE: (312) 984-9740
TELEPHONE: (312) 984-9740
TELEPHONE: 352-3856
INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 1276 base pairs
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NO. .
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ThrIleGluTyrMetAlaProGluIleLeuValArgSerGlyHisAsnArgAlaValAsp
                          AATTCTACCTGGCTGAACTTGCACTTGCTTTAGACCATCTACNTAGCCTGGGAATAATT
                                                                                                                                                           CysPheTyrLeuAlaGluIleThrLeuAlaLeuGlyHisLeuHisSerGlnGlyIleIle
                                                                                                                                                                                                AGGGGAGGAGATTTGTTTACACGCTTATCCAAAGAGGTGATGTTCACAGAAGAAGATGTC
                                                                                                                                                                                                                  SerGlyGluLeuPheThrHisLeuGluArgGluGlyIlePheLeuGluAspThrAla
                                                                                                                                                                                                                                                   GTCAAGTTGCATTATCT-TTTCAAACTGAAGGGAAGTTGTATCTTAT-TTGGATTTTCTC
                                                                                                                                                                                                                                                                      ValGluLeuAlaTyrAlaPheGlnThrGlyGlyLysLeuTyrLeuIleLeuGluCysLeu
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                                                                                                                                                                                                                                                                                                                                                                                                                           GTATTAGGGCAGGGATCATTTGGAAAGGTTTTCTTAGTTAAAAAAATCTCAGGCTCTGAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AlaAlaGlyLeuGluProValGlyHisTyrGluGluValGluLeuThrGluThrSer---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GluGlyGluGlyGluProGluLeuSerProAlaAspAlaCysProLeuAlaGluLeuArg
                                                                                   TATAGAGACTTAAAACCAGAAAATATCTT-CTTGATGAAGAAGGTCACATCAAGTTAACA
                                                                                                           TyrArgAspLeuLysProGluAspIleMetLeuSerSerGlnGlyHisIleLysLeuThr
                                                                                                                                                                                                                                                                                                             GACCGAGTTCGGACAAAAATGGAACGTGATATCTTGGTAGAGGTTAATCATCCTTTTATT
                                                                                                                                                                                                                                                                                                                                      AspThrAlaHisThrArgAlaGluArgAsnIleLeuGluSerValLysHisProPheIle
                                                                                                                                                                                                                                                                                                                                                                    GCTAGGCAGCTTTATGCCATGAAGGTATTGAAGAAGGCCACACTG-----AAAGTTCGA
                                                                                                                                                                                                                                                                                                                                                                                                                                            ValLeuGlyLysGlyGlyTyrGlyLysValPheGlnValArgLysValGlnGlyThrAsn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ------ValAsnValGlyProGluArgIleGlyProHisCysPheGluLeuLeuArg
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Matches:
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Mismatches:
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                                                                                                                                                                                                                                                                                                                                                  787 AAAAGATCGAAAAGAACAATGACTATGATTCTTAAAGCCAAAACTTGGAATGCCACAGT
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                                                                                                               GGCCTGAAGATACATTCTATTTTGATCCTGAGTTTACTGCAAAAACTCCCCAAAGATTCAC
                                                                                  roAspAspThrAlaLeuSerGluSerAlaAsnGlnAlaPheLeuGlyPheThrTyrValA
                                                                                                                                                                         TAGACTGGAATAAACTGTATAGAGA-GAAATTCATCCGCCATTTAAACCTGCAACGGGCA
                                                                                                                                                                                                                                  GATTAGGTGCAGGACCAGATGGAGTTGAAGAAATTAAAAGACAFTCATTTTTCTCAACGA
                                                                                                                                                                                                                                                   rgIleGlyGlyGlyProGlyAspAlaAlaAspValGlnArgHisProPhePheArgHisM
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Search completed: July 3, 2004, 12:18:00 Job time: 149 secs

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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Perfect score:
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Published Applications AA:*

1: /cgn2 6/ptodata/2/pubpaa/US07 PUBCOMB.pep:*

2: /cgn2 6/ptodata/2/pubpaa/PCT NEW PUB.pep:*

3: /cgn2 6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

4: /cgn2 6/ptodata/2/pubpaa/US07 NEW PUB.pep:*

5: /cgn2 6/ptodata/2/pubpaa/US07 NEW PUB.pep:*

6: /cgn2 6/ptodata/2/pubpaa/US08 NEW PUB.pep:*

6: /cgn2 6/ptodata/2/pubpaa/US08 NEW PUB.pep:*

9: /cgn2 6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

9: /cgn2 6/ptodata/2/pubpaa/US09 PUBCOMB.pep:*

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11: /cgn2 6/ptodata/2/pubpaa/US09 PUBCOMB.pep:*

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14: /cgn2 6/ptodata/2/pubpaa/US09 PUBCOMB.pep:*

15: /cgn2 6/ptodata/2/pubpaa/US09 PUBCOMB.pep:*

16: /cgn2 6/ptodata/2/pubpaa/US10B.PUBCOMB.pep:*

16: /cgn2 6/ptodata/2/pubpaa/US10B.PUBCOMB.pep:*

17: /cgn2 6/ptodata/2/pubpaa/US10 NEW PUB.pep:*

18: /cgn2 6/ptodata/2/pubpaa/US10 NEW PUB.pep:*

18: /cgn2 6/ptodata/2/pubpaa/US10 NEW PUB.pep:*

18: /cgn2 6/ptodata/2/pubpaa/US10 NEW PUB.pep:*
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2614
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(without alignments)
2798.782 Million cell updates/sec
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Listing first 45 summaries
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Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARI

No.	Score	Query Match	Query Match Length	DB	ID	Description
ב	2596	99.3	502	12	US-10-425-114-54060	Sequence 54060, A
2	1756	67.2	333	12	US-10-217-155A-16	Sequence 16, Appl
Ψ	1756	67.2	333	15	US-10-217-574-16	16,
4.	1756	67.2	333	μ÷	US-10-217-555-16	16,
G	1746	8.99	502	14	US-10-316-253-32	-
Φ	1740	66.6	525	14	US-10-316-253-34	34,
7	1428	54.6	637	ø	US-09-817-310-2	Sequence 2, Appli
8	1303	49.8	272	12	US-09-925-298-663	Sequence 663, App
9	1303	49.8	272	<b>⊨</b> 4	US-10-102-806-663	Sequence 663, App
10	1211.5	46.3	281	12	US-10-220-955-13	Sequence 13, Appl
11	1174.5	44.9	342	12	US-10-425-114-54517	545
12	955.5	36.6	544	15	US-10-108-260A-3334	Sequence 3334, Ap
13	951.5	36.4	740	15	US-10-369-493-5149	Sequence 5149, Ap
14	949.5	36.3	740	9	US-09-771-161A-265	Sequence 265, App
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Sequence 17, Appl	17,	17,	32,	32		32	Sequence 144, App	Sequence 5, Appli	Sequence 22653, A	12,		186	'n	φ,	Ø	7,					Sequence 177033,		Ñ		11	Sequence 15, Appl	Ņ	Sequence 5150, Ap	

#### ALIGNMENTS

GENERAL INFORMATION:

APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven B
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT APPLICATION NUMBER: US/10/425,114
RUMBER OF SEQ ID NOS: 73128
SEQ ID NO 54060
LENGTH: 502
TYPE: PRT Query Match 99.3%; Best Local Similarity 99.8%; %:Matches 491; Conservative ( Ś 문 Ś ; FEATURE: ; OTHER INFORMATION: Clone ID: LIB3081-101-B10\_FLI.pep US-10-425-114-54060 RESULT 1 US-10-425-114-54060 Sequence 54060, Application US/10425114 Publication No. US20040034888A1 ORGANISM: Homo sapiens 124 KDTAHTRAERNILESVKHPFIVELAYAFQTGGKLYLILECLSGGELFTHLEREGIFLEDT 183 71 11 GRRARGAGAAMAAVFDLDLETEEGSEGEGEPELSPADACPLAELRAAGLEPVGHYEEVEL 70 TETSVNVGPERIGPHCFELLRVLGKGGYGKVFQVRKVQGTNLGKIYAMKVLRKAKIVRNA 123 GRRARGAGAAMAAVFDLDLETEEGSEGEGEGEPELSPADACPLAELRAAGLEPVGHYEEVEL TETSVNVGPERIGPHCFELLRVLGKGGYGKVFQVRKVQGTNLGKIYAMKVLRKAKIVRNA 130 0 Score 2596; DB 12; Pred. No. 1.3e-178; 0; Mismatches 1; Indels Length 502; 0; Gaps

240

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APPLICANT: Barford, David
APPLICANT: Yang, Jing
APPLICANT: Yang, Jing
APPLICANT: Cron, Peter D
TITLE OF INVENTION: Kinase Crystal Structures and Materials and Methods for
TITLE OF INVENTION: Kinase Activation
FILE REPERENCE: 44236
CURRENT FILING DATE: 2002-08-14
PRIOR APPLICATION NUMBER: US/10/217,155A
CURRENT FILING DATE: 2001-08-14
PRIOR APPLICATION NUMBER: GB 0119860.5
PRIOR FILING DATE: 2001-08-14
PRIOR APPLICATION NUMBER: GB 0209985.1
PRIOR FILING DATE: 2002-05-01
NUMBER OF SEQ ID NOS: 40
SOSTWARE: Patentin Ver. 2.1
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Best Local S
Matches 331
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LENGTH: 333
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Unknown Organism: Sequence OTHER INFORMATION: uncertain
                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Unknown Organism FEATURE:
  121
                                     194
                                                                        134 NILESVKHPFIVELAYAFQTGGKLYLILECLSGGELFTHLERBGIFLEDTACFYLABITL 193
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                                                                                                                                                                                                                                                     Similarity
  ALGHLHSQGIIYRDLKPENIMLSSQGHIKLTDFGLCKESIHEGAVTHTPCGTIEYMAPBI 180
                           ALGHLHSQGIIYRDLKPENIMLSSQGHIKLTDFGLCKESIHEGAVTHTFCGTIEYMAPBI 253
                                                                                                                                                     RIGPHCFELLRVLGKGGYGKVFQVRKVQGTNLGKIYAMKVLRKAKIVRNAKDTAHTRAER
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; TITLE OF INVENTION: Kinase Crystal Structures
; TITLE OF INVENTION: 14237
; CURRENT APPLICATION NUMBER: US/10/217,574
; CURRENT FILING DATE: 2002-12-23
; PRIOR APPLICATION NUMBER: GB 0119860.5
; PRIOR FILING DATE: 2001-08-14
; PRIOR FILING DATE: 2002-05-01
; PRIOR APPLICATION NUMBER: GB 0209985.1
; PRIOR PILING DATE: 2002-07-12
; NUMBER OF SEQ ID NOS: 46
; SOFTMARE: Patentin Ver. 2.1
; SEQ ID NO 16
; SOFTMARE: Patentin Ver. 2.1
; ORGANISM: Unknown Organism
; PEATURE: ORGANISM: Unknown Organism
; OTHER INFORMATION: Description of Unknown Organism: Seq; OTHER INFORMATION: uncertain
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US-10-217-574-16
; Sequence 16, Application US/10217574
; Publication No. US20040005687A1
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Best Local Similarity
Matches 331; Conserv
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                                           RETROTEVOSPODTALSESANQAFLGETYVA
                                                                                                                    KKPLKKNPSQRIGGGPGDAADVQRHPFFRHMNWDDLLAWRVDPPFRPCLQSEEDVSQFDT
                                                                                                                                                   KKFLKRNPSORIGGGPGDAADVORHPFFRHMNWDDLLAWRVDPPFRPCLOSEEDVSOFDT
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Cron, Peter D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           67.2%; Score 1756; DB 15; ilarity 100.0%; Pred. No. 2.4e-118; Conservative 0; Mismatches 0;
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180 313

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Sequence 32, Application US/10316253
Publication No. US20030162706A1
GENERAL INFORMATION:
APPLICANT: The Procter & Gamble Company
APPLICANT: Peters, Kevin
APPLICANT: Thompson, Larry
APPLICANT: Greis, Kenneth
ITILE OF INVENTION: Angiogenesis Modulating Profile Reference: 8865M
FILE REFERENCE: 8865M
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 60/355,295
PRIOR FILING DATE: 2002-02-08
                                                                                                                                                                                                                                                                 US-10-316-253-32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Description of Unknown Organism:
; OTHER INFORMATION: uncertain
US-10-217-555-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Kinase Crystal Structures and Materials and Methods for INTEL OF INVENTION: Kinase Activation

FILE REFERENCE: 44236

CURRENT APPLICATION WIMBER: US/10/217,555

CURRENT FILING DATE: 2002-12-05

PRIOR APPLICATION NUMBER: GB 0119860.5

PRIOR APPLICATION NUMBER: GB 020985.1

PRIOR FILING DATE: 2002-05-01

NUMBER OF SEQ ID NOS: 40

SOCTWARE: Patentin Ver. 2.1

SEQ ID NO 16

LENGTH: 333

THORE THE COMMENT OF SEQ ID NOS: 40

SOCTWARE: Patentin Ver. 2.1
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Best Local Similarity
Matches 331; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Barford, David APPLICANT: Yang, Jing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Unknown Organism
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                 374
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                                                                                                                                                                                                                                                                                                                                               301 RFTRQTPVDSPDDTALSBSANQAFLGFTYVA 331
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                                                                                                                                                                                                                                                                                                                                                                                                                      KKFLKRNPSQRIGGGPGDAADVQRHPFFRHMNWDDLLAWRVDPPFRPCLQSBEDVSQFDT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hemmings, Brian A
Cron, Peter D
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; Pred. No. 2.4e-118;
0; Mismatches 0;
                                                                                                Proteins
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                      ; ORGANISM: Rattus norvegicus US-10-316-253-34
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US-10-316-253-34
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US-10-316-253-32
                                                                                                                                                                                                                                         Sequence 34, Application US/10316253
Publication No. US20030162706A1
GENERAL INFORMATION:
APPLICANT: The Procter & Gamble Company
APPLICANT: Peters, Kevin
APPLICANT: Thompson, Larry
APPLICANT: Thompson, Larry
APPLICANT: Wang, Peng
APPLICANT: Greis, Kenneth
TITLE DEFENCE: AGGENTALE Modulating Proteins
                                                                                            SOFTWARE: PatentIn version 3.1 SEQ ID NO 34
                                                                                                                              FILE REFERENCE: 8865M
CURRENT APPLICATION NUMBER: US/10/316,253
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 60/355,295
PRIOR FILING DATE: 2002-02-08
NUMBER OF SEQ ID NOS: 308
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                                                            TYPE:
                                                                              LENGTH: 525
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NUMBER OF SEQ ID NOS: 308
SOFTWARE: PatentIn version
SEQ ID NO 32
LENGTH: 502
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Best Local Similarity
                                            479 PP-SGTKKSK 487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            189 AEITLALGHLHSQGIIYRDLKPENIMLSSQGHIKLTDFGLCKESIHEGAVTHTFCGTIEY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14 MAAYFDIDLETEEGSEGEGEPELSPADACPLAE-LRAAGLEP----VGHYEEVELTETSV
                                                                                                                                                                                                              SQFDTRFTRQTFVDSFDDTALSESANQAFLGFTYVAPSVLDSIKEGFSFQPKLRSPRRLN 428
                                                                                                                                                                                                                                                                                                                                                                                                   MAPEILVRSGHNRAVDWWSLGALMYDMLFGSPPFTAENRKKTMDXIIRGKLALPPYLTFD
                                                                                                                                      SSPRVPVSPLKFSPFE----GFRPSPSLPE-PTELPLPPPLLPPPPSTT-----APLPIR
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QPNSGPYKKQ 486
                                                                                           GSPRTPVSPVKPSPGDFWGRGASASTANPQTPVEYPMETSGIEQMDVTTSGEASAPLPIR
                                                                                                                                                                                       SOFDSKFTROTPVDSFDDSTLSESANOVFLGFTYVAPSVLESVKEKFSFEPKIRSPRRFI
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US-09-817-310-2
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                                                                                                                                                       Query Match
Best Local S
Matches 279
                                                                                                                                                                                                                                                            SEQ.ID NO 2
LENGTH: 637
TYPE: PRT
ORGANISM: Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/09817310 Patent No. US20010042254A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                            APPLICANT: Stewart, Mary
APPLICANT: Kozma, Sarah
APPLICANT: Thomas, George
TITLE OF INVENTION: Drosophila Melanogaster P70 S6 Kinase
TITLE OF INVENTION UNMBER: US/09/817,310
CURRENT APPLICATION NUMBER: US/09/817,310
CURRENT FILING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: 09/230,247
PRIOR PILING DATE: 1999-04-16
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.0
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Best Local Similarity 68.0%;
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                                                                                                                                                  tch 54.6%; Score 1428; DB 9; Length 637; al Similarity 67.2%; Pred. No. 2.1e-94; 279; Conservative 47; Mismatches 81; Indels
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78 HCFELLRVLGKGGYGKVFQVRKVQGTNLGKIYAMKVLRKAKIVRNAKDTAHTRAERNILE
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                                                                DLELQDDKARDSDDBRIELDDVDLEP--ELCINLHQDTEGQETIQLCEENVNPGKIKLGP
                                                                                                           DIDLETEEGSEGEGEP-ELSPADACPLAELRAAGLEPVGHYEEVELTETSVNVGPERIGP
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; Pred. No. 5.8e-117;
51; Mismatches 83;
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PITLE OF INVENTION: Nucleic Acids, Proteins and FILE REFERENCE: PA103; CURRENT APPLICATION NUMBER: US/09/925,298; CURRENT FILING DATE: 2001-08-10; PRIOR FILING DATE: 2000-03-08; PRIOR FILING DATE: 1909-03-12; NUMBER OF SEQ ID NOS: 846; SOFTWARE: Patentin Ver: 2.0

ENGINE APPLICATION UNUMBER: S0/124,270; PRIOR FILING DATE: 1999-03-12; NUMBER OF SEQ ID NOS: 846; SOFTWARE: Patentin Ver: 2.0; SEQ ID NO 663; LENGTH: 272; Type: PAT ORGANISM: Homo sapiens; NAME/KEY: SITE; NAME/KEY: SITE; NAME/KEY: SITE; LOCATION: (29); OTHER INFORMATION: Xaa equals any of the nature of the content of the conten
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Best Local Similarity
Matches 242; Conserv
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210 APVSPLKFSPFEGFRPSPSLPBPTELPLPPLPPPLFPPPSTTAPLFIRPPSGTKKSXRGRGR 269
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Pred. No. 7.7e-86;
0; Mismatches 1;
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Sequence 13, Application US/10220955
Publication No. US20030211989A1
GENERAL INFORMATION:
APPLICANT: PLOMMAN, GREGORY D.
APPLICANT: MYTTE, DAVID
APPLICANT: MYTTE, DAVID
APPLICANT: MANNING, GERARD
APPLICANT: SUDARSANAM, SUCHA
APPLICANT: MARTINEZ, RICHAD
TITLE OF INVENTION: MOVEL HUMAN PROTEIN KINASES AND TITLE OF INVENTION: ENZYMES
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CURRENT FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: 09/925,298
PRIOR PILING DATE: 2001-08-10
PRIOR PILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05881
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR PILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 846
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 663
LENGTH: 272
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US-10-220-955-13
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; LOCATION: (29)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-10-102-806-663
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FILE REFERENCE: 038602-1401

CURRENT APPLICATION NUMBER: US/10/220,955

CURRENT FILLING DATE: 2001-03-02

NUMBER OF SEQ ID NOS: 40

SOFTMARE: Patentin Ver. 2.1

SEQ ID NO 13

LENGTH: 281
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Best Local Similarity
Matches 242; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens FEATURE:
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o. US20030054421A1
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; Pred. No. 7.7e
0; Mismatches
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7.7e-86;
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GENERAL INFORMATION:

APPLICANT: Liu, Jingdong
APPLICANT: Edou, Yihua
APPLICANT: Edou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILLE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 54517
LENGTH: 342
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
FEATURE:
OTHER INFORMATION: Clone ID: LIB4656-028-E10_Fil.pep
US-10-425-114-54517
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US-10-425-114-54517
US-10-425-114-54517, Application US/10425114
; Publication US/104034888A1
; Publication US/104034888A1
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                                                                                                                                                                                                                                                                                                                                     Query Match 44.9%; So
Best Local Similarity 70.8%; Px
Matches 225; Conservative 33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 46.3%; Score 1211.5; DB 12; Length Best Local Similarity 79.4%; Pred. No. 3.1e-79; Matches 224; Conservative 30; Mismatches 27; Indels
                                                                    133
                                                                                                                                                                                                                                           245 TIEYMAPEILVRSGHNRAVDWWSLGALMYDMLTGSPPFTAENRKKTMDKIIRGKLALPPY 304
                                                                                                        305 LTPDARDLVKKELKRNPSQRIGGGPGDAADVQRHPFFRHMNWDDLLAWRVDPPFRPCLQS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               357 PFRPCLQSEEDVSQFDTRFTRQTFVDSFDDTALSESANQAFL 398
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              181 CKLNLPPYLTQEARDLLKKLLKRNAAS-LGAGPGDAGEVQAHPFFRHINWEELLAQKVEP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        177 GIFLEDTACFYLASITLALGHLHSQGIIYRDLKPENIMLSSQGHIKLTDFGLCKBSIHEG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               117 AKIVRNAKDTAHTRABRNILESVKHPFIVELAYAPQTGGKLYLILECLSGGELFTHLERE 176
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EEDVSQFDTRFTRQTFVDSFDDTALSESANQAFLGFTYVAFSVLDSIKEGFSFQFKLRSF 424
                                                               LTOBARDILLKKLIKRNAASRIGAGPGDAGEVQAHPFFRHINWEELLARKVEPPFKPLLQS
                                                                                                                                                           TIEYMAPEILMRSGHNRAVDWWSLGALMYDMLTGAPPFTGENRKKTIDKILKCKLMLPPY
                                                                                                                                                                                                                                                                                                                                     Score 1174.5; DB 12; Length 342; Pred. No. 1.8e-76; 3; Mismatches 41; Indels 19;
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RESULT 13
US-10-369-493-5149
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US-10-108-260A-3334
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Publication No. US20040005560A1
GENERAL INFORMATION:
APPLICANT: HELIX RESEARCH INSTITUTE
APPLICANT: HELIX RESEARCH INSTITUTE
TITLE OF INVENTION: No. US20040005560A1e1 full length cDNA
FILE REFERENCE: H1-A0106
CURRENT APPLICATION NUMBER: US/10/108,260A
CURRENT FILING DATE: 2002-03-27
NUMBER OF SEQ ID NOS: 5458
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 3334
                                                                  Sequence 5149, Application US/10369493 Publication No. US20030233675A1 GENERAL INFORMATION:
  APPLICANT:
APPLICANT:
APPLICANT:
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ORGANISM: Homo sapiens
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Cao, Yongwei
Hinkle, Gregory J.
Slater, Steven C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                --ERIGPHCFELLRVLGKGGYGKVFQVRKVQGTNLGKIYAMKVLRKAKIVRNAKDTAHTR
                                                                                                                                                                                                                                                                                     PRVPVSPL 438
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                                           Yongwei
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; Pred. No. 1.8e-60;
74; Mismatches 105;
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US-10-369-493-5149
                      ; ORGANISM: Homo sapiens
US-09-771-161A-265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-771-161A-265
                                                                                         PRIOR APPLICATION NUMBER: 135619
PRIOR FILING DATE: 2000-04-12
NUMBER OF SEQ ID NOS: 273
SOFTWARE: PatentIn version 3.0
SEQ ID NO 265
LENGTH: 740
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 265, Application US/09771161A Patent No. US20020110811A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 5149
LENGTH: 740
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                APPLICANT: LEVINE, et al.
TITLE OF INVENTION: VARIANTS OF PROTEIN KINASES
FILE REFERENCE: 802620-2005.1
CURRENT APPLICATION NUMBER: US/09/771,161A
CURRENT FILING DATE: 2001-01-26
                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 09/724,676
PRIOR FILING DATE: 2000-11-8
PRIOR APPLICATION NUMBER: 136776
PRIOR FILING DATE: 2000-06-15
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APPLICANT: Chen, Xianfeng
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10(52052)B
FILE REFERENCE: 38-10(52052)B
FILE REFERENCE: 38-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
CURRENT FILING DATE: 2003-02-28
                                                                         TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            103 HISHPFİVKLHYAPQTEGKLYLİLDFİRĞGDİFTRİSKEVMFTEDDVKFYLAELTLALEH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    138 SVKHPFIVELAYAFQTGGKLYLILECLSGGELFTHLEREGIFLEDTACFYLAEITLALGH
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47.9%; Pred. No. 5.1e-60;
active 69; Mismatches 105;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -SETEIDIGDVRKCGBKADP
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CURRENT APPLICATION NUMBER: US/09/771,161A
CURRENT FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 09/724,676
PRIOR APPLICATION NUMBER: 09/724,676
PRIOR FILING DATE: 2000-11-28
PRIOR PILING DATE: 2000-06-15
PRIOR PILING DATE: 2000-06-15
PRIOR APPLICATION NUMBER: 135619
PRIOR FILING DATE: 2000-04-12
NUMBER OF SEQ ID NOS: 273
SOFTWARE: PATENTING DATE: 2000-04-12
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; Sequence 266, Application US/09771161A
; Patent No. US20020110811A1
; GENERAL INFORMATION:
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Best Local Simi
Matches 185;
                                                                                                                                                                                                                                                                                                                                                                                Matches 185;
                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: LEVINB, et al.
TITLE OF INVENTION: VARIANTS OF PROTEIN KINASES
FILE REFERENCE: 802620-2005.1
131 VKLHYAFQTEGKLYLILDFLRGGDLFTRLSKEVMFTEEDVKFYLAELALALDHLHSLGII 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   371
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                                     VELAYAFQTGGKLYLILECLSGGELFTHLEREGIFLEDTACFYLAEITLALGHLHSQGII 204
                                                                                                                                                        VIGKGGYGKVFQVRKVQGTNLGKIYAMKVLRKAKIVRNAKDTAHTRAERNILESVKHPFI 144
                                                                                                                      VLGQGSFGKVFLVKKISGSDARQLYAMKVLKKATL--KVRDRVRTKMERDILVEVNHPFI
                                                                                                                                                                                                                                                                                                                EGEGEPELSPADACPLAELRAAGLEPVGHYEEVELTETS----VNVGPERIGPHCFELLR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LGAGPDGVEEIKRHSFFSTIDWNKLYRREIHPFFKPATGRPEDTFYFDPEFTAKTPKDSP 370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DDTALSESANQAFLGFTYVA 404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           #WSLGALMYDMLTGSPPFTAENRKKTMDKIIRGKLALPPYLTPDARDLYKKFLKRNPSQR 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           YRDLKPENIMLSSQGHIKLTDFGLCKESIHEGAVTHTFCGTIEYMADEILVRSGHNRAVD 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VKLHYAFQTBGKLYLILDFLRGGDLFTRLSKEVMFTBEDVKFYLAELALALDHLHSLGII 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VLGKGGYGKVFQVRKVQGTNLGKIYAMKVLKXAKIYRNAKDTAHTRAERNILESVKHPFI 144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EGEGEPELSPADACPLAELRAAGLEPVGHYEEVELTETS----VNVGPERIGPHCFELLR 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -GIPPSANAHQLFRGFSFVA 389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IGGGPGDAADVQRHPFFRHMNWDDLLAWRVDPPFRPCLQSEEDVSQFDTRFTRQTPVDSP 384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            YRDLKPENILLDEEGHIKLTDFGLSKESIDHEKKAYSFCGTVEYMAPEVVNRRGHTQSAD 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VELAYAFQTGGKLYLILECLSGGELFTHLEREGIFLEDTACFYLAEITLALGHLHSQGII 204
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                                                                                                                                                                                                                                                                                                                                                                      36.3%; Score 949.5; DB 9;
48.7%; Pred. No. 7.1e-60;
ative 65; Mismatches 105;
                                                                                                                                                                                                                                                   ------QTEEVSIKEIAITHHVKEGHEKADPSQFELLK
                                                                                                                                                                                                                                                                                                                                                                            Indels 25;
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                                                                                                                                                                                                                                             191 YRDIKPENILLDESGHIKLTDFGLSKESIDHEKKAYSFCGTVEYMAPEVVNRRGHTQSAD
                                                                               311 LGAGEDGVEEIKRHSFFSTIDWNKLYRREIHFFFKPATGRPEDTFYFDFEFTAKTFKDSP 370
                                                                                                                                                               251 WWSPGVLMFEMLTGTLPFQGKDRKETMTWILKAKLGMPQFLSPEAQSLLRMLFKRNPANR 310
                                                                                                                                                                                                                                                                                           205
                                     DDTALSESANQAFLGFTYVA 404
                                                                                                                                                                                     WWSLGALMYDMLTGSPPFTAENRKKTWDKIIRGKLALPPYLTPDARDLVKKFLKRNPSQR 324
                                                                                                                                                                                                                                                                   YRDIKPENIMLSSOGHIKLTDFGLCKESIHEGAVTHTFCGTIEYMAPEILVRSGHNRAVD
-GIPPSANAHQLFRGFSFVA 389
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Search completed: June 30, 2004, 17:20:18
Job time: 51 secs

US-10-425-114-18480 US-09-968-433-10

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Command line parameters:

-MODEL-frame+ p2n.model -DEV=xlh
-Q-/Ggn2 1/USPTO_spool/US0762258/runat_30062004_165610_13661/app_query.fasta_1.647
-Q-/Ggn2 1/USPTO_spool/US0762258/runat_30062004_165610_13661/app_query.fasta_1.647
-DB=Published_Applications_NA -QFMT=fastap -SUPFTX=rnpb -KINMATCH=0.1
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXIEN=2000D000000 -USER=US09762258 GCGN 1 487 @runat 30062004 165610 13661
-NCQU=5 -1CQU=3 -NO_MANA -LARGBOUGHEY -NGE_SCORES=0 -MAIT -DSPBLOCK=100
-LONGLOG -DEV_TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5
-FGAPOD=6 -FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Minimum DB seq length: Maximum DB seq length:
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Xgapop 10.0 , Xgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BLOSUM62
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2614
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/cgn2_6/ptodata/2/pubpna/USO6_PUBCOMB.seq:*
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ALIGNMENTS

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) ORGANISM: HOMO SAPLENS ; FEATURE: OTHER THEORMATION: Clone ID: LIRYOR1-101-R10 FLI	; TYPE: DNA	; LENGTH: 1742	; SEQ ID NO 18480	; NUMBER OF SEQ ID NOS: 73128	; CURRENT FILING DATE: 2003-04-28	; CURRENT APPLICATION NUMBER: US/10/425,114	; FILE REFERENCE: 38-21(53313)B	प्र TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement	; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With	; APPLICANT: Cao, Yongwei	; APPLICANT: Tabaska, Jack B	; APPLICANT: Screen, Steven E	; APPLICANT: Kovalic, David K.	; APPLICANT: Zhou, Yihua	; APPLICANT: Liu, Jingdong	; GENERAL INFORMATION:	; Publication No. US20040034888A1	; Sequence 18480, Application US/10425114	US-10-425-114-18480	RESULT 1

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

Result No. Score Query Match Length DB IJ

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                                                                                                                                  AlaCysPheTyrLeuAlaGluIleThrLeuAlaLeuGlyHisLeuHisSerGlnGlyIle
TyrLeuThrProAspAlaArgAspLeuValLysLysPheLeuLysArgAsnProSerGln
                                          AlaGluAsnArgLysLysThrMetAspLysIleIleArgGlyLysLeuAlaLeuProPro
                                                                                               AspTrpTrpSerLeuGlyAlaLeuMetTyrAspMetLeuThrGlySerProPheThr
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                                                                               GACTGGTGGAGCCTGGAGCCCTGATGTACGACATGCTCACTGGATCGCCGCCCTTCACC
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SEQ ID NO 10

LENGTH: 1637

TYPE: DNA

ORGANISM: Homo sapiens

PATURE:
HAME/KEY: misc feature
OTHER INFORMATION: Incyte ID No. US20030073162A1 1869688

NAME/KEY: unsure
LOCATION: 3, 13, 22, 44, 69, 162, 1220, 1426, 1443, 1458, 1465, 1486, 1480, 1490, 100CATION: 1517, 1522, 1524-1525, 1533, 1553, 1573, 1584, 1605, 1624, 1631, 1634-1

OTHER INFORMATION: a, t, c, g, or other
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 10, Applica Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                      APPLICANT: Lal, Preeti
APPLICANT: Au-Young, Janice
APPLICANT: Reddy, Roopa
APPLICANT: Murry, Lynn E.
APPLICANT: Murry, Lynn E.
APPLICANT: Mathur, Preete
TITLS OF INVENTION: SIGNAL PEPTIDE-CONTAINING PROTEINS
FILE REFERENCE: PC-0051 CIP
CURRENT APPLICATION UMBER: US/09/968,433
CURRENT FILING DATE: 2001-10-01
NUMBER OF SEQ ID NOS: 79
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                          yProGlyAspAlaAlaAspValGlnArgHisProPhePheArgHisMetAsnTrpAspAs 348
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                                                  Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                Alignment Scores:
                                                                                                                                                                                    US-09-954-456-1137
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CURRENT FILING DATE: 2001-09-18
PRIOR APPLICATION NUMBER: US/60/23,617
PRIOR FILING DATE: 2000-09-18
PRIOR PRIOR PAPPLICATION NUMBER: US/60/234,052
PRIOR FILING DATE: 2000-09-20
PRIOR PPLICATION NUMBER: US/60/234,923
PRIOR PPLICATION NUMBER: US/60/234,923
PRIOR PILING DATE: 2000-09-25
PRIOR APPLICATION NUMBER: US/60/235,134
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR APPLICATION NUMBER: US/60/235,638
PRIOR PILING DATE: 2000-09-26
PRIOR APPLICATION NUMBER: US/60/235,638
PRIOR APPLICATION NUMBER: US/60/235,711
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LENGTH: 2346
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NUMBER OF SEQ ID NOS: 2276
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                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/60/235,720 FILING DATE: 2000-09-27 APPLICATION NUMBER: US/60/235,840 FILING DATE: 2000-09-27 APPLICATION NUMBER: US/60/235,863 FILING DATE: 2000-09-27
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TrpAspAspLeuLeuAlaTrpArgValAspProProPheArgProCysLeuGlnSerGlu
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; Publication No. US20030083284A1
; GENERAL INFORMATION:
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APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0245
CURRENT FILLING DATE: 2001-08-01
RUMBER: OF SEQ ID NOS: 49
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TGGGAAGAACTTCTGGCTCGAAAGGTGGAGCCCCCCTTTAAACCTCTGTTGCAATCTGAA
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Best Local Similarity:
Query Match:
DB:
                                                                                              Score:
 US-09-762-258-2 (1-495)
                                                                                                                            Alignment Scores
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US-10-388-360-321
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SOFTWARE: FastSEQ for Windows
SEQ ID NO 321
LENGTH: 2346
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                                                                                                                                                                                                                                                                  APPLICANT: GENOMIC HEALTH
APPLICANT: Baker, Joffre B.
APPLICANT: Baker, Joffre B.
APPLICANT: Cronin, Maureen T.
APPLICANT: Kiefer, Michael C.
APPLICANT: Shak, Steve
APPLICANT: Malker, Michael Graham
TITLE OF INVENTION: GENE EXPRESSION PROFILING IN BIOPSIED TUMOR TISSUES
TILE REFERENCE: 39740-0001US
CURRENT APPLICATION NUMBER: US/10/388,360
CURRENT FILING DATE: 2003-03-12
PRIOR APPLICATION NUMBER: US 60/412,049
PRIOR APPLICATION NUMBER: US 60/364,890
PRIOR APPLICATION NUMBER: US 60/364,890
PRIOR PILING DATE: 2002-03-13
PRIOR FILING DATE: 2002-03-13
                                                                                                                                                                           ORGANISM: Homo
                                                                                                                                                                                            TYPE: DNA
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x US-10-388-360-321 (1-2346)
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                             Length:
Matches:
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                                   GGAGCTGGTCCTGGGGACGCTGGAGAAGTTCAAGCTCATCCATTCTTTAGACACATTAAC
                                                                                                      ACACAAGAAGCCAGAGATCTGCTTAAAAAAGCTGCTGAAAAAGAAATGCTGCTTCTCGTCTG
                                                                                                                                                                                               AsnArgLysLysThrMetAspLysIleIleArgGlyLysLeuAlaLeuProProTyrLeu
                                                                                                                                                                                                                                                TGGAGTTTGGGAGCATTAATGTATGACATGCTGACTGGAGCACCCCCATTCACTGGGGAG
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                                                                                                                                                                                                                                                                                                                                         IleGluTyrMetAlaProGluIleLeuValArgSerGlyHisAsnArgAlaValAspTrp
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                                                        GlyGlyGlyProGlyAspAlaAlaAspValGlnArgHisProPhePheArgHisMetAsn
                                                                                                                                      ThrProAspAlaArgAspLeuValLysLysPheLeuLysArgAsnProSerGlnArgIle
                                                                                                                                                                            NATAGAAAGAAAACAATTGACAAAATCCTCAAATGTAAACTCAATTTGCCTCCCTACCTC
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                                                                                                                   ; LENGTH: 1800
; TYPE: DNA
; ORGANISM: Rattus norvegicus
; FEATURB:
; FEATURB:
; LOCATION: (134)...(1642)
; OTHER INFORMATION:
US-10-316-253-31
Percent Similarity:
Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                              APPLICANT: Wang, Feng
APPLICANT: Greis, Kenneth
TITLE OF INVENTION: Angiogenesis Modulating Pr
FILE REFERENCE: 8865M
CURRENT APPLICATION NUMBER: US/10/316,253
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 60/355,295
PRIOR FILING DATE: 2002-02-08
NUMBER OF SEQ ID NOS: 308
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Sequence 31, Application US/10316253
Publication No. US20030162706A1
GENERAL INFORMATION:
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                                                                       Pred. No.:
                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1 SEQ ID NO 31
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APPLICANT: Peters, Kevin
APPLICANT: Thompson, Larry
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                                                                                                   AlaArgAspLeuValLysLysPheLeuLysArgAspProSerGlnArgIleGlyGlyGly
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TYPE: DNA
CRCANUISM: Rattus norvegicus
FRATURB:
NAME/KEY: CDS
LOCATION: (22)...(1599)
OTHER INFORMATION:
US-10-316-253-33
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Publication No. US20030162706A1

GENERAL INFORMATION:

APPLICANT: The Procter & Gamble Company

APPLICANT: Thompson, Larry

APPLICANT: Wang, Feng

APPLICANT: Wang, Feng

APPLICANT: Greis, Kenneth

TITLE OF INVENTION: Angiogenesis Modulating Pr

FILE REFERENCE: 8865M
                                                                                   US-09-762-258-2 (1-495) x US-10-316-253-33 (1-2287)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/316,253
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 60/355,295
PRIOR FILING DATE: 2002-02-08
NUMBER OF SEQ ID NOS: 308
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ATGAGGCGACGACGGAGGCGGGACGGCTTTTACCCAGCGCCTGACTTCCGACACAGGGGAA 81
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                                                                       TrpAspAspLeuLeuAlaTrpArgValAspProProPheArgProCysLeuGlnSerGlu
                                                                                                                                   GGAGCTGGCCCTGGGGATGCTGGAGAAGTCCAAGCGCATCCATTTTTTAGACACATTAAC
                                                                                                                                                        GlyGlyProGlyAspAlaAlaAspValGlnArgHisProPhePheArgHisMetAsn 345
                                                                                                                                                                                                                 ACACAAGAAGCTCGAGATCTGCTTAAAAAGCTGCTGAAGAGAAATGCTGCTTGTCTTT
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APPLICANT: ELASHOFP, Michael

ITITE OF INVENTION: Cardiotoxin Molecular Toxicology Modeling

FILE REFERENCE: 44921-5090US

CURRENT APPLICATION NUMBER: US(10/191,803)

CURRENT APPLICATION NUMBER: US(60/303,819)

PRIOR APPLICATION NUMBER: US(60/303,819)

PRIOR FILING DATE: 2001-07-10

PRIOR APPLICATION NUMBER: US(60/305,623)

PRIOR APPLICATION NUMBER: US(60/305,623)

PRIOR APPLICATION NUMBER: US(60/305,351)

PRIOR APPLICATION NUMBER: US(60/307,611)

PRIOR APPLICATION NUMBER: US(60/377,611)

PRIOR APPLICATION NUMBER: US(60/377,611)

PRIOR FILING DATE: 2002-04-03

PRIOR APPLICATION NUMBER: US(60/377,611)

PRIOR FILING DATE: 2002-05-06

NUMBER OF SEQ ID NOS: 1140

SOPTWARE: Patentin Ver. 2.1

SEQ ID NO 230

LENGTH: 2287

TYPE: DNA

ORGANISM: Rattus norvegicus

FEATURE:

OTHER INFORMATION: Genbank Accession No. US20040014040A1 NM_0319
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APPLICANT: PORTER, Mark
APPLICANT: JOHNSON, Kory
APPLICANT: HIGGS, Brandon
APPLICANT: CASTLE, Arthur
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    ATGAGGCGACGACGGAGGCGGGACGGCTTTTACCCAGCGCCTGACTTCCGACACAGGGAA
                                       MetAlaArgGlyArgArgAlaArgGlyAlaGlyAlaAla----
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APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Screen, Steven E
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack B
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REPERENCE: 38-21 (53313) B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEO, ID NOS: 73128
SEO, ID NO 26874
LENGTH: 2362
TYPE: DNA
OPCANISM: Homo sapiens
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Publication No. US20040034888A1
GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
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Best Local Similarity:
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OTHER INFORMATION: Clot
US-10-425-114-26874
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                                                                                                                                                  ATGAGGCGACGAAGGAGGCGGGACGCCTTTTACCCAGCCCCGGACTTCCGAGACAGGGAA
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Matches:
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Publication No. US20020039764A1

GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA103
CURRENT APPLICATION NUMBER: US/09/925,298
CURRENT APPLICATION NUMBER: US/09/925,298
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05881
PRIOR APPLICATION NUMBER: FCT/US00/05881
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 846
SQFTWARE: PatentIN Ver. 2.0
SEQ ID NO 245
LENGTH: 1197
TYPE: DNA
ORGANISM: Homo Sapiens
FEATURE:
NAME/KEY: misc feature
LOCATION: (318)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (1193)
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Best Local Similarity:
Query Match:
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Sequence 245, Application US/10102806
Publication No. US20030054421A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins
FILE REFERENCE: FALOSPICE:
CURRENT APPLICATION NUMBER: US/10/102,806
CURRENT FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: 09/925,298
PRIOR TILING DATE: 2001-09-10
PRIOR FILING DATE: 2001-03-10
PRIOR FILING DATE: 2001-03-08
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PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 846
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 245
LENGTH: 1197
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ORCANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
LOCATION: (218)
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OTHER INFORMATION: n equals a,t,
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                                              rProAspAspTbrAlaLeuSerGluSerAlaAsnGlnAlaPheLeuGlyPheThrTyrVa
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Patent No. US20010042254A1
GENERAL INFORMATION:
APPLICANT: Stewart, Mary
APPLICANT: Kozma, Sarah
APPLICANT: Thomas, George
TITLE OF INVENTION: Drosophila Melanogaster
FILE REFERENCE: 4.20971/A
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LENGTH: 2556
TYPE: DNA
ORGANISM: Drosophila
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CURRENT FILLING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: 09/230,247
PRIOR FILING DATE: 1999-04-16
NUMBER OF SEQ ID NOS: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.0
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                   GlnGlyThrAsnLeuGlyLysIleTyrAlaMetLysValLeuArgLysAlaLysIleVal
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SEQ ID NO 58
LENGTH: 1058
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APPLICANT: Au-Y
APPLICANT: Redd
APPLICANT: Murr
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APPLICANT: Reddy, Roopa
APPLICANT: Mathur, Preete
FITLE OF INVENTION: SIGNAL PEPTIDE-CONTAINING PROTEINS
FILE REFERENCE: PC-0051 CIP
CURRENT APPLICATION NUMBER: US/09/968,433
CURRENT FILING DATE: 2001-10-01
NUMBER OF SEQ ID NOS: 79
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ORGANISM: Mus musculus
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OTHER INFORMATION:
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                                              NAME/KEY: unsure
                                                                                                               TYPE: DNA
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Sequence 126, Application US/10305720
Publication No. US20040010136A1
GEMERAL INFORMATION:
APPLICANT: Au-Young, Janice K.; Seilhamer, Jeffrey J.
TITLE OF INVENTION: Composition for the Detection of Signaling Pathway Gene Express
FILE REFERENCE: PA-0002-1 COM
FILE REFERENCE: PA-0002-1 COM
CURRENT APPLICATION NUMBER: US/10/305,720
CURRENT FILING DATE: 2002-11-26
PRIOR APPLICATION NUMBER: 09/016,434
PRIOR FILING DATE: 1998-01-30
NUMBER OF SEQ ID NOS: 1490
NUMBER OF SEQ ID NOS: 1490
                                                                                                                                                                                                                  SOFTWARE: PERL Program
SEQ ID NO 126
LENGTH: 1607
                                                                                              ORGANISM: Homo sapiens FEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Incy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GlyThrIleGluTyrMetAlaProGluIle-LeuValArgSerGlyHisAsnArgAlaVa
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Qy 336 nArgHisProPhePheArgHisMetAsnTrpAspAspLeuLeuAlaTrpArgValAspPr 356	Qy 316 eLeuLysArgAsnProSerGlnArgIleGlyGlyGlyGlyFroGlyAspAlaAlaAspValGl 336	Qy 296 gGlyLysLeuAlaLeuProProTyrLeuThrProAspAlaArgAspLeuValLysLysPh 316	Qy 276 uThrGlySerProProPheThrAlaGluAsnArgLysLysThrMetAspLysIleIleAr 296	O nArgAlaValAspTrpTrpSerLeuGlyAlaLeuWetTyrAspMetLe	Qy 254	Qy 245 rIleGluTyrMetAlaProGluIle 253	Qy 229CysLysGluSerIleHisGluGlyAla-ValThrHisThrPheCysGlyTh 245	Qy 218 GlnGlyHisIleLysLeuThrAspPheGlyLeu 228 :::   ::    :        Db 605 CCCAACCAGGTCTCCCCCCTGAAGATCTGTGACTTCGAGCTGGGCAGCGGCATCAAACTC 664	Qy 198 LeuHisSerGlnGlyIleIleTyrArgAspLeuLysProGluAsnIleMetLeuSerSer 217	Qy 178 ilePheLeuGluAspThrAlaCysPheTyrLeuAlaGluIleThrLeuAlaLeuGlyHis 197	Qy 158 TyrLeuIleLeuGluCysLeuSerGlyGlyGluLeuPheThrHisLeuGluArgGluGly 177	92 8	Qy 118 LysIleValArgAsnAlaLysAspThrAlaHisThrArgAlaGluArgAsnIleLeuGlu 137 :::::       ::	Qy 106GlyLysIleTyrAlaMetLysValLeuArgLysAla 117 :::       ::: ::::: Db 272 ATCACCAGCCAGGAGTACNCCGTCAANATCATTGNTTTTTTNCAGGCCACATTCGGAGC 331	Qy 86 LeuGlyLysGlyGlyTyrGlyLysValPheGlnValArgLysValGlnGlyThrAsnLeu 105	16 Gaps: 8-2 (1-495) x US-10-305-720-126 (1-1607)	Alignment Scores:  Pred. No.:  5.13e-131
OY 157 LeuTyrLeuIleLeuGluCysLeuSerGlyGlyGluLeuPheThrHisLeuGluArgGlu 176	Qy 137 GluSerValLysHisProPheIleValGluLeuAlaTyrAlaPheGlnThrGlyGlyLys 156	Qy 117 AlaLysIleValArgAsnālaLysAspThrAlaHisThrArgālaGluĀrgĀsnīleLeu 136	13 Gaps: -762-258-2 (1-495) x US-10-220-955-1 (1-843)	: 1211.50 In Similarity: 90.07% Local Similarity: 79.43% Match: 46.35%	S-10-220-955-1  S-10-220-955-1  Lignment Scores: 1.12e-113 Length:	O 1 : 843 DNA SM: Homo sapiens	CURRENT CURRENT NUMBER SOFTWAR	APPLICANT: MARTINEZ, RICARDO TITLE OF INVENTION: NOVEL HUMAN PROTEIN KINASES AND PROTEIN KINASE-LIKE TITLE OF INVENTION: ENCYMES FILE REFERENCE: 038602-1401		10-22	1498 CATCCGTCCCCCTCAGGGACCAAGAAGTCCAAGAGGGCCGTGGGCGTCCAGGGCGT	1438 1438		1318 CTTCCAGCCCAGCCTCACCAGGCGCCTCAACAGTAGCCCCGGGTCCCCGTCAG	396 1258 416		Qy 356 oProPheArgProCysLeuGlnSerGluGluAspValSerGlnPheAspThrArgPheTh 376

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Search completed: July Job time : 711 secs
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                                                                          ArgHisProPhePheArgHisMetAsnTrpAspAspLeuLeuAlaTrpArgValAspPro 356
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                                                                                                                                                                                                                                                                                                        SerGlyHisAsnArgAlaValAspTrpTrpSerLeuGlyAlaLeuMetTyrAspMetLeu 276
                                                                                                                                                                                                                                                                                                                                              AlaValThrHisThrPheCysGlyThrIleGluTyrMetAlaProGluIleLeuValArg
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                                                                                                                                                                                                                                                                                                                                                                                                                         HisLeuHisSerGlnGlyIleIleTyrArgAspLeuLysProGluAsnIleMetLeuSer 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                              GlyIlePheLeuGluAspThrAlaCysPheTyrLeuAlaGluIleThrLeuAlaLeuGly 196
                                                                                                                                                                                         LeuLysArgAsnProSerGlnArgIleGlyGlyGlyProGlyAspAlaAlaAspValGln 336
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Scoring table:
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Perfect score:
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                                     3163042 segs, 2412103800 residues
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Copyright (c) 1993 - 2004 Compugen Ltd.

    2004, 09:58:27; Search time 842 Seconds
(without alignments)
    10404.704 Million cell updates/sec
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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries seg length: seg length: 20000000000

Database 10: 11: 12: 13: 14: 15: 16: 18: Published Applications NA:\*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

\_6/ptodata/2/pubpna/US60\_PUBCOMB.seq:\*

# SUMMARIES

Result No.	Score	Query Match	Length DB	DB	ID	Description
1	1706.2	94.0	1742	13	US-10-425-114-18480	Sequence 18480, A
ń	1158.6	63.8	1197	13	US-09-925-298-245	Sequence 245, App
<del>د</del>	1158.6	63.8	1197	15	US-10-102-806-245	
4	1104.4	60.8	1637	10	US-09-968-433-10	10, A
Ųι	720	39.6	1607	16	US-10-305-720-126	•
φ	672.6	37.0	1058	10	US-09-968-433-58	58,
7	628.8	34.6	1800	15	US-10-316-253-31	31,
8	628.2	34.6	2287	15	US-10-316-253-33	33,
ý.	628.2	34.6	2287	16	US-10-191-803-230	
10	619.8	34.1	2346	9	US-09-954-456-1137	Sequence 1137, Ap
11	619.8	34.1	2346	10	US-09-920-677-3	Sequence 3, Appli
12	619.8	34.1	2346	13	US-10-388-360-321	Sequence 321, App
13	593.6	32.7	669	16	US-10-305-720-118	Sequence 118, App
14	0 77	2	2260		TIC_10_42E_114_2E074	Segmence 26874 A

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253.8	253.8	٠	259	259	ហ	٠		260.8			260.8			293.2	297.2	297.2	305.8	331.2	332.4	334	337	337	371	392.6	398	418.8	418.8	455.6	٠
14.0	14.0	14.1	14.3	14.3	14.3	14.4	14.4	14.4	14.4	14.4	14.4	14.4	14.4	16.1	16.4	16.4	16.8	18.2	18.3	18.4	18.6	18.6	20.4	21.6	21.9	23.1	23.1	25.1	0.0
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17	15	17	17	15	15	17	17	16	φ	15	15	15	10	15	13	13	13	φ	φ	16	15	16	φ	10	10	10	10	13	5
US-10-641-643-1004	US-10-324-985A-3	-10-713-678-3	-10-713-67	24-	-10-324-9	US-10-713-678-1	US-10-641-643-1206	US-10-388-263-329	US-09-970-000-3	US-10-072-036-138	US-10-072-036-70	US-10-198-846-9984	US-09-955-999-43	US-10-112-286-1	US-10-172-118-972	US-10-342-887-972	US-10-240-315-2	US-09-880-107-181	US-09-880-107-2146	US-10-159-563-155	US-10-240-965-120	US-10-108-260A-891	US-09-925-300-418	US-09-918-995-29197	US-09-968-433-59	US-09-873-319-168	US-09-960-706-276	US-10-220-955-1	00-00-000-00'
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## ALIGNMENTS

GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Lou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven B
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei APPLICANT: Tabaska, Jack B
APPLICANT: Tabaska, Jack B
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: 15313)B
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 18480
LENCTH: 1742
TYPE: DNA
ORGANISM: Homo sapiens US-10-425-114-1848D Sequence 18480, Application US/10425114 Publication No. US20040034888A1

밁 S 밁 Ś ; FEATURE: ; OTHER INFORMATION: Clone ID: LIB3081-101-B10\_FLI US-10-425-114-18480 Query Match Best Local Similarity Matches 1708; Conserv 146 ACGGAGGAAGGCAGCGAGGGCGAGGGCCAGAGCTCAGCCCCGCGGACGCATGTCCC 86 GGCCGACGGGCCCGGCGGGCCGCCATGGCGGCCGTCTTTGATTTGGATTTGGAG 32 GGCCGACGGGCCCGCGGGGCCGCCCCCATGGCGGCCGTGTTTGATTTGGATTTGGAG Conservative 94.0%; Score 1706.2; Pred. No. 0; Mismatches DB 13; Length 1742; ب --Indels <u>,</u> Gaps 205 151 91 145

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CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05881
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 846
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 245
LENGTH: 1197
TYPE: DNA
ORGANIEM: Homo Bapiens
                                                                                                                                                    FEATURE:

NAME/KEY: misc feature

LOCATION: (218)

LOCATION: 118)

OTHER INFORMATION: n equals a,t,g, or NAME/KEY: misc feature

LOCATION: (1193)

OTHER INFORMATION: n equals a,t,g, or US-09-925-298-245
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publication No. US20020039764A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins
FILE REFERENCE: PA103
                                                                                 Query Match 63.8
Best Local Similarity 99.3
Matches 1180; Conservative
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99.3%;
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Pred. No. 6.6e-302;
3; Mismatches 3;
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LENGTH: 1197
TYPE: DNA
ORGANISM: Homo Bapiens
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LOCATION: (218)
OTHER INFORMATION: n equals a
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CURRENT FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: 09/925,298
PRIOR FILING DATE: 2001-08-10
PRIOR FILING DATE: 2001-08-01
PRIOR APPLICATION NUMBER: 05/120/05881
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 846
SOPTWARE: Patentin Ver. 2.0
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Best Local
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GENERAL INFORMATION
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TITLE OF INVENTION: Nucleic Acids, Proteins
FILE REFERENCE: PA103P1C1
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APPLICANT: Lal, Preeti
APPLICANT: Au-Young, Janice
APPLICANT: Reddy, Roopa
APPLICANT: Reddy, Roopa
APPLICANT: Mathur, Preete
TITLS OF INVENTION: SIGNAL PEPTIDE-CONTAINING PROTEINS
FILS REFERENCE: PC-0051 CIP
CURRENT APPLICATION NUMBER: US/09/968,433
CURRENT FILING DATE: 2001-10-01
NUMBER OF SEQ ID NOS: 79
SOFTWARE: PERL PROGRAM
SEQ ID NO 10
LENGTH: 1637
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TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: Incyte ID No. US20030073162A1 1869688
NAME/KEY: unsure
LOCATION: 3, 13, 22, 44, 69, 162, 1220, 1426, 1443, 1458, 1465, 1486, 1488,
LOCATION: 1517, 1522, 1524-1525, 1533, 1553, 1573, 1584, 1605, 1624, 1631, 1
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Best Local Similarity
Matches 1196; Conserv
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RESULT 5
US-10-305-720-126
Sequence 126, Application US/10305720
Publication No. US20040010136A1
GENERAL INFORMATION:
APPLICANT: AU-YOUNG, Janice K.; Seilhamer, Jeffrey J.
TITLE OF INVENTION: Composition for the Detection of S
FILE REFERENCE: PA-0002-1 CON
CURRENT APPLICATION NUMBER: US/10/305,720
CURRENT APPLICATION NUMBER: 09/016,434
PRIOR APPLICATION NUMBER: 09/016,434
PRIOR FILING DATE: 1998-01-30
NUMBER OF SEQ ID NOS: 1490
SOFTWARE: PERL Program
SEQ ID NO 126
LENGTH: 1507
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OTHER INFORMATION:
US-10-305-720-126
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ORGANISM: Homo sapiens
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OTHER INFORMATION:
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                   GGACGTGAGCCAGTTTGATACCCGCTTCACACGGCAGACGCCGGTGGACAGTCCTGATGA 1233
   GGACGTGAGCCAGTTTGATACCCGCTTCACACGCCAGACGCCGGTGGACAGTCCTGATGA
                                                                                                                                           GGGTGGCCCAGGGGATGCTGATGTGCAGAGACATCCCTTTTTCCGGCACATGAATTG
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                                                             GGACGACCTTCTGGCCTGGCGTGTGGACCCCCCTTTCAGGCCCTGTCTGCAGTCAGAGGA
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Pred. No. 1.1e-183;
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APPLICANT: Lal, Preeti
APPLICANT: Nurry, Janice
APPLICANT: Murry, Lynn B.
APPLICANT: Murry, Lynn B.
APPLICANT: Mathur, Preete
TITLE OF INVENTION: SIGNAL PEPTIDE-CONTAINING PROTEINS
FILE REFERENCE: PC-0051 CIP
CURRENT APPLICATION NUMBER: US/09/968,433
CURRENT APPLICATION NUMBER: US/09/968,433
CURRENT FILING DATE: 2001-10-01
NUMBER OF SEQ ID NOS: 79
SOFTWARE: PERL Program
SEQ ID NO 58
LENGTH: 1058
TYPE: DNA
ORGANISM: Mus musculus
FEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID NO. US20030073162A1 037196_Mm.1
US-09-968-433-58
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Best Local Similarity
Matches 745; Conserv
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ACTGAGACCAGCGTGAACGTTGGCCCAGAGCGCATCGGGCCCCACTGCTTTTGAGCTGCTG
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                                                CTTGGCGAATTAAGGGCTGCTGGCCTGGAGACAGTGGGACACTATGAAGAAGTAGAGCTG
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                                                                                                                                                                                                                                  Score 672.6; DB 10
Pred. No. 5.8e-171;
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Query Match 34.6%; Score 628.8; DB 15; Length 1800; Best Local Similarity 71.1%; Pred. No. 4.1e-159;	NAME/KEY: CDS ; LOCATION: (134)(1642) ; CTHER INFORMATION: US-10-316-253-31	INGTH: TPE: DN TGANISM		316,253	; APPLICANT: Thompson, Larry ; APPLICANT: Wang, Feng ; APPLICANT: Wang, Kenneth ; TITLE OF INVENTION: Anglogenesis Modulating Proteins	; Publication No. US20030162706A1 ; GENERAL INFORMATION: ; APPLICANT: The Procter & Gamble Company ; APPLICANT: Peters, Kevin	RESULT 7 US-10-316-253-31 ; Sequence 31, Application US/10316253	Qy 865 GGACTGGTGGAGCCTGGGGGCCCTGATGTACGACATGCTCACTGGATCGCC 915	Oy 806 GGCACCATTGAGTACATGGCCCCTGAGA-TTCTGGTGCGCAGTGGCCACAACCGGGCTGT 864	Qy 746 ACCGACTTIGGACTCTGCAAGGAGTCTATCCATGAGGGGGCGCTCACTCA	QY 686 ATCTACCGGGACCTCAAGCCCGAGAACATCATGCTCAGCAGCCAGGGCCACATCAAACTG 745	Qy 626 GCCTGCTTCTACCTGGCTGAGATCACGCTGGGCCATGTCCCACTCCCAGGGCATC 685	Qy 566 CTCAGTGGTGGCGAGCTCTTCACGCATCTGGAGCGAGAGGGCATCTTCCTGGAAGATACG 625	QY 506 ATTGTGGAACTGGCCTATGCCTTCCAGACTGGTGGCAAACTCTACCTCATCCTTGAGTGC 565	QY 446 AAGGACACAGACACACACGGCTGAGCGGAACATTCTAGAGTCAGTGAAGCACCCCTTT 505	Qy 386 AACITGGGCAAAATATATGCCATGAAAGTCCTAAGGAAGGCCAAAATTGTGCGCAATGCC 445	Qy 326 CGTGTGCTGGGCAAGGGGGGCTATGGCAAGGTGTTCCAGGTGCGAAAGGTGCAAGGCACC 385	Db 228 ACAGAGAGCAGCGTGAACCTGGGTCCTGAGCGCATCGGGCCCCACTGCTTTGAGCTACTG 287
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1281 AGGTCTTTCTGGGTTTTACATATGTGGCTCCATCTGTACTTGAAAGTGTGAAAGAAA	TCACACGGCAGACGCCGGTGGACAGTCCTGATGACACACGCCTCAGCGAGAGAGTGCCAACC	1140 ACCCCCTTTCAGGCCCTGTCTGCAGTCAGAGGAGGAGGTGAGCCAGTTTGATACCCGCT 1199	1080 TGCAGAGACATCCCTTTTTCCGGCACATGAATTGGGACCTTCTGGCCTGGCGTGGG 1139	1020 AGTTTCTGAAACGGAATCCCAGCCAGCGGATTGGGGGTGGCCCAGGGGATGCTGGTGATG 1079	960 TCAGGGGCAAGCTGGCACTGCCCCCTACCTCACCCCAGATGCCCGGGACCTTGTCAAAA 1019 	900 TGCTCACTGGATGGCGCCCTTTACCGCAGAGAACCGGAAGAAACCATGGATAAGATCA 959	TGCGCAGTGGCCACAACCGGGCTGTGGACTGGTGGAGCCTGGGGGGCCCTGATGTACGACA	780 AGGGCGCCGTCACTCACACCTTCTGCGGCACCATTGAGTACATGGCCCCTGAGATTCTGG 839	720 TCAGCAGCCAGGGCCACATCAAACTGACCGACTTTGGACTCTGCAAGGAGTCTATCCATG 779	660 TGGGCCATCTCCACTCCCAGGGCATCATCTACCGGGACCTCAAGCCCGAGAACATCATGC 719	600 GAGAGGGGATCTTCCTGGAAGATACGGCCTGCTTCTACCTGGCTGAGATCACGCTGGCCC 659	540 GCAAACTCTACCTCATCCTTGAGTGCCTCAGTGGTGGGGGGGCTCTTCACGCATCTGGAGC 599	480 TTCTAGAGTCAGTGAAGCACCCCTTTATTGTGGAACTGGCCTATGCCTTCCAGACTGGTG 539	420 GGAAGGCCAAAATTGTGCGCAATGCCAAGGACACAGCACACACA	360 TCCAGGTGCGAAAGGTGCAAGGCACCAACTTGGGCAAAATATATGCCATGAAAGTCCTAA 419	300 TCGGGCCCCACTGCTTTGAGCTGCTGCGTGTGCTGGGCAAGGGGGGGCTATGGCAAGGTGT 359	240 TGGGACACTATGAAGAGGTGGAGCTGAGACCAGCGTGAACGTTGGCCCAGAGCGCA 299	nes 831; Conservative 0; Mismatches 337; Indels 0; Gaps $0;$

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; LOCATION: (22)..(1599)
; OTHER INFORMATION:
US-10-316-253-33
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US-10-316-253-33
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Matches 830
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SEQ ID NO 33
LENGTH: 2287
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Publication No. US20030162706A1
GENERAL INFORMATION:
APPLICANT: The Procter & Gamble Company
APPLICANT: Peters, Kevin
APPLICANT: Thompson, Larry
APPLICANT: Wang, Feng
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TITLE OF INVENTION: Angiogenesis Modulating
FILE REFERENCE: 8865M
CURRENT APPLICATION NUMBER: US/10/316,253
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 60/355,295
PRIOR FILING DATE: 2002-02-08
PRIOR FILING DATE: 2002-03-08
NUMBER OF SEQ ID NOS: 308
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ORGANISM: Rattus
FEATURE:
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Sequence 230, Application US/10191803
| Publication No. US20040014040A1
| GENERAL INFORMATION: APPLICANT: MENDRICK, Donna | APPLICANT: MENDRICK, Donna | APPLICANT: HIGGS, Brandon | APPLICANT: HIGGS, Brandon | APPLICANT: LIGGS, Brandon | APPLICANT: ELASHOFF, Michael | TITLE OF INVENTION: Cardiotoxin Molecular Toxicology Modeling | CURRENT FILING DATE: 2002-07-10 | PRIOR APPLICATION NUMBER: US 60/303,819 | PRIOR APPLICATION NUMBER: US 60/303,819 | PRIOR APPLICATION NUMBER: US 60/305,623 | PRIOR APPLICATION NUMBER: US 60/305,623 | PRIOR APPLICATION NUMBER: US 60/369,351 | PRIOR APPLICATION NUMBER: US 60/369,351 | PRIOR APPLICATION NUMBER: US 60/369,351 | PRIOR APPLICATION NUMBER: US 60/369,351 | PRIOR APPLICATION NUMBER: US 60/369,351 | PRIOR APPLICATION NUMBER: US 60/377,611 | PRIOR PILING DATE: 2002-04-03 | PRIOR APPLICATION NUMBER: US 60/377,611 | PRIOR FILING DATE: 2002-05-06
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; FEATURE:
; OTHER INFORMATION: Genbank Accession No. US20040014040A1 NM_031985
US-10-191-803-230
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; SOFTWARE: PatentIn V
; SEQ ID NO 230
; LENGTH: 2287
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Best Local Similarity 71.1
Matches 830; Conservative
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TYPE: DNA
ORGANISM: Rattus norvegicus
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Pred. No. 6.3e-159;
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PRIOR FILING DATE: 2000-09-27;
NUMBER OF SEQ ID NOS: 2276-
; SOPTWARE: Patentin version 3.0;
SEQ ID NO 1137;
LENGTH: 2346;
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CURRENT FILING DATE: 2001-09-18

PRIOR APPLICATION NUMBER: US/60/233,617

PRIOR FILING DATE: 2000-09-18

PRIOR FILING DATE: 2000-09-20

PRIOR APPLICATION NUMBER: US/60/234,052

PRIOR FILING DATE: 2000-09-25

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PRIOR APPLICATION NUMBER: US/60/235,134

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RESULT 11

US-09-920-677-3

Sequence 3, Application US/09920677

Publication No. US20030083284A1

GENERAL INFORMATION:
APPLICANT: Earctt P. Monia
APPLICANT: Lex M. Cowsert

TITLE OF INVENTION: ANVISENSE MODULATION OF P70

FILE REFERENCE: RTS-0245

CURRENT APPLICATION NUMBER: US/09/920,677

CURRENT APPLICATION NUMBER: US/09/920,677

CURRENT FILING DATE: 2001-08-01

NUMBER OF SEQ ID NOS: 49

SEQ ID NO 3

LENGTH: 2346

TYPE: DNA

CORGANISM: Homo sapiens

FEATURE:
NAME/KEY: CDS

LOCATION: (28)...(1605)

US-09-920-677-3
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Best Local Similarity 70.5%;
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TGCTCACTGGATCGCCGCCCTTTACCGCAGAGAACCCAGGAAGAAACCATGGATAAGATCA
                                                      TGAGAAGTGGCCACAATCGTGCTGGATTGGTGGAGTTTGGGAGCATTAATGTATGACA
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                                                                                    TGCGCAGTGGCCACAACCGGGCTGTGGACTGGTGGAGCCTTGGGGGGCCCTGATGTACGACA
                                                                                                                ATGGAACAGTCACACACATTTTGTGGAACAATAGAATACATGGCCCCTGAAATCTTGA
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Pred. No. 1.2e-156;
0; Mismatches 344;
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APPLICANT: WALKER, Michael Graham
TITIE OF INVENTION: GENE EXPRESSION PROFILING IN
FILE REFERENCE: 39740-0001US
CURRENT APPLICATION NUMBER: US/10/388,360
CURRENT FILING DATE: 2003-03-12
PRIOR APPLICATION NUMBER: US 60/412,049
PRIOR FILING DATE: 2002-09-18
PRIOR PRIOR DATE: 2002-09-18
PRIOR FILING DATE: 2002-09-13
INUMBER OF SEQ ID NOS: 384
SECTIMARE: FastSEQ for Windows Version 4.0
LENGTH: 2346
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Publication No. US203022555
GENERAL INFORMATION:
APPLICANT: GENOMIC HEALTH
APPLICANT: Baker, Joffre E
APPLICANT: Cronin, Maureer
APPLICANT: Kiefer, Michael
APPLICANT: Shak, Steve
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US-10-388-360-321
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Best Local Similarity
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ORGANISM: Homo sapiens
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  TCCAGGTGCGAAAGGTGCAAGGCACCAACTTGGGCAAAATATATGCCATGAAAGTCCTAA 419
                                              TCAGACCAGAATGTTTTGAGCTACTTCGGGTACTTGGTAAAGGGGGGCTATGGAAAGGTTT
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Kiefer, Michael C.
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; OTHER INFORMATION: a, t,
US-10-305-720-118
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TITLE OF INVENTION: Composition for the Detection of Signaling Pathway
FILE REPERENCE: PA-0002-1 CON
CURRENT APPLICATION NUMBER: US/10/305,720
CURRENT FILING DATE: 2002-11-26
PRIOR APPLICATION NUMBER: 09/016,434
PRIOR PRILING DATE: 1998-01-30
PRIOR FILING DATE: 1998-01-30
NUMBER OF SEQ ID NOS: 1490
SOFTWARE: PERL Program
SEQ ID NO 118
LENGTH: 669
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Best Local Similarity
Matches 646; Conserv
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Pred. No. 1e-149;
0; Mismatches 18;
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APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
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APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated |
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION UNMEER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 26874
LENGTH: 2362
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Best Local Similarity
Matches 886; Conserv
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Pred. No. 9.4e-134;
0; Mismatches 422;
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Sequence 1, Application US/09817310
PALEAI NO. US20010042254A1
GENERAL INFORMATION:
APPLICANT: Stewart, Mary
APPLICANT: Kozwa, Sarah
APPLICANT: Thomas, George
TITLE OF INVENTION: Drosophila Melanogaster P70
FILE REFERENCE: 4-20971/A
CURRENT APPLICATION NUMBER: US/09/817,310
CURRENT FILING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: 09/230,247
PRIOR FILING DATE: 1999-04-16
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